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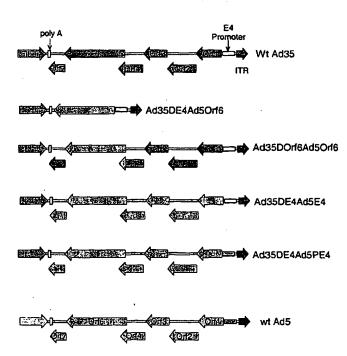
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[Continued on next page]

#### (54) Title: METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY



(57) Abstract: Various methods propagating and rescuing multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line are Typically, replication-defective disclosed. adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The disclosed methods offer the ability to propagate vectors derived from multiple adenoviral serotypes in a single production cell line which expresses E1 proteins from a single serotype. Propagation in this manner is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the expressed E1 of the cell line and the heterologous E4 of the replication-defective adenoviral vectors enables their propagation and rescue. The invention bypasses a need in the art to customize specific cell lines to the serotype or subgroup of the adenoviral vector being propagated and enables one to easily and rapidly develop alternative adenoviral serotypes as gene delivery vectors for use as vaccines or as a critical component in gene therapy.

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#### TITLE OF THE INVENTION

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METHODS FOR PROPAGATING ADENOVIRUS AND VIRUS PRODUCED THEREBY

## CROSS-REFERENCE TO RELATED APPLICATIONS

The present application claims the benefit of application serial nos. 60/458,825, filed March 28, 2003; 60/455,312, filed March 17, 2003; 60/455,234, filed March 17, 2003; and 60/405,182, filed August 22, 2002.

#### FIELD OF THE INVENTION

The present invention concerns various methods to propagate and rescue multiple serotypes of replication-defective adenovirus in a single adenoviral E1-complementing cell line. Typically, replication-defective adenovirus vectors propagate only in cell lines which express E1 proteins of the same serotype or subgroup as the vector. The methods disclosed herein offer the ability to propagate vectors derived from multiple serotypes in a single cell line expressing E1 proteins from a single serotype. Such propagation of a wide range of vectors in one cell line is accomplished by providing all or a portion of an E4 region in cis within the genome of the replication-defective adenovirus. The added E4 region or portion thereof is cloned from a virus of the same or highly similar serotype as that of the E1 gene product(s) of the complementing cell line. Interaction between the E1 gene products of the cell line and the heterologous E4 gene products of the replication-defective adenoviral vector enables the propagation and rescue of the recombinant replication-defective adenovirus vectors. The invention, therefore, bypasses an existing need in the art to customize complementing cell lines to the specific serotype or subgroup of the adenoviral vector being propagated or, alternatively, to have to transfect a cell line with an E4 region and then regulate the expression in trans of the E4 region within the E1 complementing cell line.

## BACKGROUND OF THE INVENTION

Beginning with the first human adenoviruses (Ads) isolated over four decades ago (Rowe et al., Proc. Soc. Exp. Biol. Med., 84:570-579, 1953), over 100 distinct serotypes of adenovirus have been isolated which infect various mammalian species, 51 of which are of human origin (Straus, Adenovirus infections in humans. In The Adenoviruses. 451-498, 1984; Hierholzer et al., J. Infect. Dis., 158: 804-813, 1988; Schnurr and Dondero, Intervirology., 36: 79-83, 1993; Jong et al., J Clin Microbiol., 37:3940-3945:1999). The human serotypes have been categorised into six subgenera (A-F) based on a number of biological, chemical, immunological and structural criteria; criteria which include hemagglutination properties of rat

and rhesus monkey erythrocytes, DNA homology, restriction enzyme cleavage patterns, percentage of G+C content and oncogenicity (Straus, Adenovirus infections in humans. In *The Adenoviruses*. 451-498, 1984; Horwitz, Adenoviridae and their replication, *In Virology*: 1679-172, 1990).

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Deletion of an essential E1 region common to the various adenovirus serotypes has enabled the use of adenovirus vectors as gene transfer vectors for vaccine and gene therapy purposes. Resultant replication-defective vectors are propagated in cell lines that provide the deleted E1 gene products in trans. Supplementation of the essential E1 gene products in trans in this manner works well when the E1 gene products are from the same or a highly similar serotype. As such, E1-deleted group C serotypes (Ad1, Ad2, Ad5 and Ad6) grow well in 293 or PER.C6 cells which contain and express the Ad5 E1 region. In contrast, E1-deleted serotypes other than group C, for example those from subgroups A, B, D, E, and F (e.g., Ad3, Ad4, and Ad7 to Ad51), do not replicate efficiently in 293 or PER.C6 cells. The Ad5 E1 sequences in 293 and PER.C6 cells do not fully complement the replication of these alternative serotypes. This presents a challenge due to the fact that the most characterized and studied complementing cell lines available for growth and propagation of adenovirus are based on E1 sequence from adenovirus serotype 5.

This inability to fully complement the replication of serotypes other than group C adenovirus in Ad5 E1 complementing cell lines has been attributed to the inability of Ad5 (group C) E1b 55K gene product to functionally interact with the E4 gene products of non-group C serotypes. While the interaction is conserved within members of the same subgroup, it is not well conserved between subgroups.

Hence, cell lines expressing both Ad5 E1 and ORF6 were generated and proved useful in complementing alternative adenovirus serotypes; see, e.g., Abrahamsen et al., 1997 J. Virol. 8946-8951. Such incorporation of E4 (or ORF6) into Ad 5 complementing cell lines as was done in Abrahamsen et al., supra, is known.

U.S. Patent No. 5,849,561 discloses complementation of an E1-deleted non-group C adenovirus vector in an Ad5-E1 complementing cell line which also expresses portions of the Ad5-E4 gene.

U.S. Patent No. 6,127,175, issued to Vigne, *et al.*, discloses a stably transfected mammalian cell line which expresses a portion of the E4 region of adenovirus, preferably ORF6 or ORF6/7. Such a cell line is useful for complementation of recombinant Ad genomes deficient in the E4 region.

European Application EP 1 054 064 A1 discloses recombinant, replication deficient adenovirus 35 (Ad35) vectors and cell lines which complement *in trans* the growth of

these vectors. A cell line which expresses Ad5E1A and E2A genes (PER.C6) was shown to complement an Ad35-E1 deleted vector upon co-expression of Ad35-E1B proteins.

U.S. Patent No. 6,270,996, issued to Wilson, et al., discloses E1/E4 deleted adenovirus vectors and E1/E4(ORF6) cell lines which complement in trans virus growth without resulting in cell toxicity.

U.S. Patent No. 6,202,060, issued to Mehtali, et al., discloses adenoviral vectors wherein portions of the early genes are under control of an inducible promoter. The '060 patent also discloses complementing cell lines which may be used in tandem with these Ad vectors.

The generation of serotype-specific cell lines providing a complementing serotype-specific E1 gene product(s) in trans is known as well.

Although Ad5-based vectors have been used extensively in a number of gene therapy trials, there may be limitations on the use of Ad5 and other group C adenoviral vectors due to preexisting immunity in the general population due to natural infection. Ad5 and other group C members tend to be among the most seroprevalent serotypes. Immunity to existing vectors may develop as a result of exposure to the vector during treatment. These types of preexisting or developed immunity to seroprevalent gene delivery vectors may limit the effectiveness of gene therapy or vaccination efforts. Alternative adenovirus serotypes, thus, constitute very important targets in the pursuit of gene delivery systems capable of evading the host immune response.

There remains both a practical and commercial need for an adenovirus-based vaccine and/or gene therapy delivery system which allows for the production of multiple serotype recombinant adenovirus vectors in a single source complementing mammalian cell line. The present invention addresses and overcomes this deficiency in the art by disclosing novel methods for propagating multiple serotype recombinant Ad vectors in a single complementing cell line where the required serotype-specific sequences are provided *in cis*.

#### SUMMARY OF THE INVENTION

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The present invention relates to an enhanced means for propagating replication-defective adenovirus in an E1-complementing cell line(s) where the E1 gene product(s) being expressed is not native to the adenovirus being propagated. The method is based on Applicants' finding that supply, in cis, of a nucleic acid sequence encoding all or a portion of a heterologous adenoviral E4 region which is native to a virus of the same or highly similar serotype as the E1 gene product(s) of the complementing cell line enables the growth of adenoviral vectors of varying serotype in any single complementing cell line, despite the fact the cell line is not customized for the particular serotype of vector being propagated. This is of particular

importance given that existing and settled adenoviral E1-complementing cell lines (such as PER.C6™ and 293) are based on one of the most prominent adenovirus serotypes (Ad5) and are not suited for the large-scale propagation and rescue of alternative serotypes.

The basic steps involved in the propagation of adenoviral vectors in accordance with the methods of the instant invention are as follows: First, all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding at least open reading frame 6 (ORF6) is inserted into a replication-defective adenoviral vector. By "heterologous", Applicants mean that the nucleic acid sequence is not native to the viral vector being propagated, i.e., not normally present within a virus of the same or highly similar serotype. As will be described, the adenoviral E4 region or portion thereof can be either a nucleic acid sequence encoding ORF 6 or any larger portion of the E4 region, and includes nucleic acid comprising the complete E4 region with E4 promoter. The region into which the nucleic acid is incorporated is not limited, i.e., the insertion can be made into the complete E4 region with E4 promoter or into a smaller portion narrowing into the ORF6 region. Alternatively, the heterologous E4 region or portion thereof can be inserted into different areas of the genome such as the E1 or E3 regions. Further, the native E4 region or portion thereof can be deleted and replaced, or left intact. This is not deemed a critical element of the instant invention. What is a critical element is that the heterologous E4 region or portion thereof being inserted is native to a virus of the same or highly similar serotype as the E1 gene product(s) expressed by the complementing cell line.

Following the modification of the adenoviral vector of interest, the recombinant adenovirus is then introduced into an adenoviral E1-complementing cell line and allowed to propagate. The adenovirus is subsequently harvested and rescued from the complementing cell line.

The resultant virus can be studied and used in various gene therapy and vaccine efforts. The virus, therefore, forms an important aspect of the instant invention.

## BRIEF DESCRIPTION OF THE DRAWINGS

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FIGURE 1 illustrates a transcription map for adenovirus serotype 5. The linear genome is divided into 100 map units as well as into r- and l- strands which designate the direction of transcription. Early transcription units are designated with an E and are active prior to viral DNA replication. Late transcription units are designated with and L and are active primarily after DNA replication. Promoters are represented as brackets and polyadenylation sites as arrowheads. The tripartite leader is designated 1, 2, and 3.

FIGURES 2A-1 through 2A-10 illustrate the nucleic acid sequence for the wild-type adenovirus 35 (SEQ ID NO: 1) utilized in the Examples.

FIGURE 3 illustrates the homologous recombination scheme utilized to recover pAd35 $\Delta$ E1.

FIGURE 4 illustrates the various configurations of the E4 regions (or portions) within the alternative serotype recombinants.

FIGURE 5 illustrates the homologous recombination scheme utilized to recover pAd35 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

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FIGURE 6 illustrates the nucleic acid sequence encoding the gag expression cassette (SEQ ID NO: 2). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of HIV-1 gag; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 7 illustrates the nucleic acid sequence encoding the SEAP expression cassette (SEQ ID NO: 3). The various regions of the figure are as follows: (1) a first underlined segment of nucleic acid sequence encoding the immediate early gene promoter region from human cytomegalovirus; (2) a first segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; (3) a region in caps which contains the coding sequence of the human placental SEAP gene; (4) a second segment of lowercase letters which is not underlined, which segment of DNA contains a convenient restriction enzyme site; and (5) a second underlined segment, this segment containing nucleic acid sequence encoding a bovine growth hormone polyadenylation signal sequence.

FIGURE 8 illustrates *in vivo* expression of SEAP in C3H/HeN mice using 10<sup>10</sup> vp doses of Ad35 vectors. This experiment was designed to address any effects of E3 deletion. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Shown are geometric means for each cohort of 5 mice.

vp doses of Ad35 vectors. This experiment was designed to address any effects of Ad5 sequence insertion into the Ad35 genome. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10^10 vp and 10^9 vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURES 10A-B illustrate in vivo SEAP expression using MRKAd5-based (A) and Ad35ΔE1ΔE4Ad5Orf6-based (B) vector in rhesus macaques. Shown are the serum antigen

levels for individual monkeys following a single intramuscular (i.m.) injection of 10<sup>11</sup> vp MRKAd5SEAP (filled circles), 10<sup>9</sup> vp MRKAd5SEAP (open boxes) or 10<sup>11</sup> vp Ad35ΔE1SEAPΔE4Ad5Orf6.

FIGURE 11 illustrates in vivo SEAP expression in African green monkeys using Ad5- and Ad35-based vectors. Shown are the antigen levels for each animal in serum samples collected two days after the treatment.

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FIGURE 12 illustrates the homologous recombination scheme utilized to recover pAd24 $\Delta$ E1.

FIGURE 13 illustrates the homologous recombination scheme utilized to recover pAd24ΔE1Ad5Orf6.

FIGURE 14 illustrates the configuration of E4 regions in the Ad24 recombinants generated.

FIGURE 15 illustrates the growth kinetics of the Ad24-based vectors in PER.C6 cells.

FIGURES 16A-1 through 16A-10 illustrate the nucleic acid sequence for wild-type adenovirus serotype 24 (SEQ ID NO: 5). The ATCC product number for Ad24 is VR-259.

FIGURE 17 illustrates, in tabular format, gag-specific T cell responses in monkeys immunized with MRKAd5-HIVgag and Ad24 HIV vectors. Shown are the numbers of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag peptide pool. The pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

FIGURE 18 illustrates, in tabular format, the characterization of the gag-specific T cells in monkeys immunized with 10^11 vp of MRKAd5-HIV1gag and Ad24ΔE1gagΔOrf6Ad5Orf6. Shown are the percentages of CD3+ T cells that are either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.03%).

FIGURE 19 illustrates individual anti-p24 titers (in mMU/mL) in macaques immunized with gag-expressing adenovirus vectors.

FIGURE 20 illustrates *in vivo* expression of SEAP in C3H/HeN mice using 10^10 vp doses of Ad24 vectors. The vectors were injected intramuscularly and the levels of SEAP expression were determined from the serum samples. Two extra cohorts received 10^10 vp and 10^9 vp of Ad5 vector. Shown are geometric means for each cohort of 5 mice.

FIGURE 21 illustrates in vivo SEAP expression using MRKAd5 and Ad24 vectors in rhesus macaques. Shown are the geometric means of the SEAP levels for cohorts of 3 monkeys. In bars are the standard errors of the geometric means.

FIGURE 22 illustrates a homologous recombination scheme to be utilized to recover pAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

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FIGURE 23 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad5/Ad6 prime-Ad24 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 or 3 doses of the priming vaccine; c: 3 wks prior to boost; d: 4 wks after boost; e: ND, not determined.

FIGURE 24 illustrates, in tabular format, the percentages of CD3<sup>+</sup> T lymphocytes that are gag-specific CD8<sup>+</sup> cells or gag-specific CD4<sup>+</sup> cells determined after the Ad24 Boost Immunization (wk 60). Numbers reflect the percentages of circulating CD3+ lymphocytes that are either gag-specific CD4+ or gag-specific CD8+ cells. Mock values (equal to or less than 0.01%) have been subtracted.

FIGURE 25 illustrates gag-specific T cell responses in rhesus macaques immunized following a heterologous Ad 24 prime-Ad5 boost regimen. a: Mock, no peptide: gag, 20-mer peptide pool encompassing entire gag sequence; b: Peak response after 2 doses of the priming vaccine; c: Wk 24; d: 4 wks after boost; e: ND, not determined.

FIGURE 26 illustrates the homologous recombination scheme utilized to recover pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

FIGURE 27 illustrates the homologous recombination scheme utilized to recover pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

FIGURES 28A-1 to 28A-9 illustrate a nucleic acid sequence for wild-type adenovirus serotype 34 (SEQ ID NO: 12). The ATCC product number for Ad34 is VR-716.

FIGURE 29 illustrates the time course of SEAP expression using MRKAd5 and Ad34 vectors in rhesus macaques. Data represent cohort geometric means.

FIGURE 30 illustrates, in tabular format, T cell responses induced using MRKAd5 and Ad34 vectors expressing HIV-1 gag. Data are expressed in numbers of spot-forming cells per million PBMC (SFC/10^6 PBMC). "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 31 illustrates, in tabular format, the levels of CD4+ and CD8+ Gagspecific T cells in Ad34-immunized macaques at week 12. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 32 illustrates, in tabular format, T cell responses induced using a heterologous Ad34 prime/Ad35 boost regimen in macaques. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

FIGURE 33 illustrates, in tabular format, the levels of CD4+ and CD8+ Gag-specific T cells in Ad34 primed/Ad35 boosted macaques at week 28. "a" refers to a 20-mer peptide pool with 10-aa overlap and encompassing the entire HIV-1 CAM1 gag.

## DETAILED DESCRIPTION OF THE INVENTION

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The present invention details an efficient strategy for the propagation and rescue of alternative adenoviral serotypes utilizing available adenovirus production cell lines, nullifying the need to customize available cell lines for a specific serotype of interest. This is enabled by the incorporation of a critical E4 region into the adenovirus to be propagated.

The critical E4 region in the instant invention comprises, in the minimum, nucleic acid sequence encoding E4 ORF6 and can comprise the entire region of E4, inclusive of the promoter region. An important characteristic of the imported E4 region is that it is native to a virus of the same or highly similar serotype as the E1 gene product(s) (particularly E1B 55K) of the E1-complementing cell line, but heterologous to (i.e., non-native to a virus of the same serotype as) the adenoviral vector being propagated. As will be detailed below, the heterologous E4 region or portion thereof can be varied and can be inserted into the vector backbone at numerous locations.

The heterologous E4 region or portion thereof can, for instance, be a nucleic acid sequence encoding the entire open reading frame of the non-native E4. This segment of nucleic acid sequence can, in turn, be incorporated into the "native" entire E4 open reading frame of the recipient virus. In such an embodiment, the promoter native to the adenoviral vector would drive the expression of the non-native E4 region within the recombinant replication-defective adenoviral vector. Alternatively, the nucleic acid sequence encoding the entire open reading frame can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

In another embodiment, the heterologous E4 region comprises a nucleic acid sequence encoding the entire open reading frame of E4 and includes a non-native E4 promoter. In this type of embodiment, the E4 region can be inserted into the location of the combined native E4 and E4 promoter region. The non-native E4 region in this embodiment would be driven by expression of the non-native E4 promoter. Alternatively, the nucleic acid sequence encoding the entire open reading frame and the non-native E4 promoter can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

An alternative and further embodiment exists wherein the heterologous E4 region or portion thereof comprises nucleic acid sequence encoding a partial E4 region comprising ORF6 (one aspect of which is a region solely encoding ORF6). In this particular aspect of the invention, the heterologous non-native E4 protein can, in certain embodiments, replace the non-native ORF6 region or the entire E4-encoding region of the native virus. In the latter situation, the promoter driving expression of the non-native ORF6 can either be the native E4 promoter or a heterologous, non-native promoter operatively linked to the non-native ORF6, while in the latter, the expression of the non-native ORF6 would generally be driven by the native E4 promoter. Alternatively, the nucleic acid sequence encoding a partial E4 region comprising ORF 6 can be inserted into a different region of the adenoviral vector genome, such as for example the E1 or E3 regions. In this latter embodiment, the native E4 region or portion thereof can be deleted or left intact.

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As one of skill in the art can appreciate, there are various ways in which one can envision the supply of a heterologous E4 nucleic acid sequence in cis to an adenoviral vector and thereby enable its growth based on Applicants' novel findings herein. Moreover, as one of skill in the art can appreciate, either native or non-native promoters can be utilized to drive expression of the heterologous E4 region or portion thereof.

Adenovirus pre-plasmids (plasmids comprising the genome of the replication-defective adenovirus with desired deletions and insertions) can be generated by homologous recombination using adenovirus backbones and an appropriate shuttle vector (designed to target-in specific deletions and incorporate desired restriction sites into the resultant plasmid). Shuttle vectors of use in this process can be generated using general methods widely understood and appreciated in the art, e.g., PCR of the adenoviral terminal ends taking into account the desired deletions, and the sequential cloning of the respective segments into an appropriate cloning plasmid. The adenoviral pre-plasmid can then be digested and transfected into the complementing cell line via calcium phosphate co-precipitation or other suitable means. Virus replication and amplification then occurs, a phenomenon made evident by notable cytopathic effect. Infected cells and media are then harvested after viral replication is complete (generally, 7-10 days post-transfection).

It is to be noted that various alternative adenoviral serotypes can be developed in accordance with the disclosed methods and, particularly, alternative adenoviral serotype vectors that were previously unable to be propagated or very inefficiently propagated utilizing existing adenoviral production cell lines based on subgroup C complementing E1 sequence. The various adenoviral vectors that can be developed in accordance with the instant methods include adenoviral vectors of subgroups A-F (for instance, serotypes of subgroups A, B (e.g., serotypes

11, 14, 16, 21, 34 and 35), C (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F.

In preferred embodiments, the various non-group C family members can be developed with heterologous E4 supplied from a subgroup C member such as adenovirus serotype 5. Particular embodiments of the instant invention utilize a development scheme wherein the heterologous E4 protein is derived from a wildtype adenovirus serotype 5 sequence; see, e.g., a viral sequence which has been deposited with the American Type Culture Collection ("ATCC") under ATCC Deposit No. VR-5 (for which a transcription map can be found in Figure 1). A particular example of this type of embodiment is wherein an adenovirus of subgroup B (or any non-C subgroup) comprising heterologous E4 proteins in cis from Ad5 is propagated in Ad5 E1-complementing cell lines, for instance, PER.C6<sup>TM</sup> or 293. Applicants have, in fact, successfully propagated E1- serotypes 10, 24, 34, and 35 via use of this particular embodiment.

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One of skill in the art can readily identify alternative adenovirus serotypes (e.g., alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C, (e.g., serotypes 2 and 5), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) for the supply of the heterologous E4 protein. As long as the heterologous E4 region (or portion thereof comprising ORF6) of the vector is native to a virus of the same or highly similar serotype as the E1 region of the complementing cell line, the methods of the instant invention are widely applicable to the propagation and rescue of adenovirus of all serotypes. In light of the present disclosure, one can readily envision, for instance, how a complementing cell line based on a non-subgroup C adenovirus (e.g., the Ad35 cell line of EP 1 054 064 A1) can be utilized to propagate a virus of an adenoviral vector of subgroup C (e.g., adenovirus serotype 5) provided that the appropriate nucleic acid sequence encoding an E4 protein provided in cis is native to a virus of the same or highly similar serotype as that of the E1 expressed by the complementing cell line (i.e., an Ad35 E4 protein).

Complementing cell lines of use in the instant invention are available in the art and are not limited to any specific type. The critical feature, again, is that the heterologous segment of E4-encoding nucleic acid sequence provided *in cis* to the replication-defective vector being propagated be native to a virus of the same or highly similar serotype as the E1 expressed by the complementing cell line. One aspect of the instant invention employs E1-complementing cell lines wherein the expressed E1 is of serotype 5; *e.g.*, PER.C6<sup>TM</sup> and 293 cell lines. Both these cell lines express the adenoviral E1 gene product. PER.C6<sup>TM</sup> is described in Fallaux *et al.*, 1998 *Human Gene Therapy* 9:1909-1917, hereby incorporated by reference. 293 cell lines are described in Graham *et al.*, 1977 *J. Gen. Virol.* 36:59-72, hereby incorporated by reference.

Another aspect of the instant invention are the adenoviral vectors of any serotype falling with adenoviral subgroups A, B, C, D, E and F (for instance, alternative serotypes of subgroups A, B (e.g., serotypes 11, 14, 16, 21, 34 and 35), C (e.g., serotype 2), D (e.g., serotypes 24, 26 and 36), E (e.g., serotype 4) and F) which are modified to contain a non-native E4-encoding nucleic acid sequence in cis which comprises, in whole or in part, nucleic acid sequence encoding open reading frame 6 (ORF6). Virus in accordance with this description can be propagated in accordance with the above-described methods and rescued using any suitable means known in the art.

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Another aspect of the instant invention is a vector in accordance with the instant invention which comprises a heterologous passenger gene in addition to that of the heterologous E4 nucleic acid sequence. In specific embodiments, the passenger gene encodes an antigen.

As one of ordinary skill in the art will appreciate, the instant methods are not limited by the heterologous gene that can be incorporated. The instant invention relates generally to a means by which to propagate multiple serotypes of adenovirus in a single complementing cell line and the recombinant virus that make the process possible. In preferred embodiments, the passenger gene is incorporated into the E1 deletion. In alternatively preferred embodiments, the passenger gene is inserted in an E3-deleted region. The position of the passenger gene, as one of ordinary skill in the art will appreciate, can be varied according to the specific complementing cell utilized and the specific deletions present within the replication-defective adenovirus genome.

In specific embodiments the passenger gene can encode an HIV-1 antigen, and in more preferred embodiments selected from the group consisting of genes encoding HIV-1 gag, pol, nef and env. In alternative embodiments, the passenger gene can be a reporter gene, such as secreted alkaline phosphatase (SEAP).

The passenger gene preferably exists in the form of an expression cassette. A gene expression cassette preferably comprises (a) a nucleic acid sequence encoding a protein of interest; (b) a promoter operatively linked to the nucleic acid sequence encoding the protein; and (c) a transcription termination sequence. The transcriptional promoter of the adenoviral vector is preferably recognized by an eukaryotic RNA polymerase. In a preferred embodiment, the promoter is a "strong" or "efficient" promoter. An example of a strong promoter is the immediate early human cytomegalovirus promoter (Chapman *et al.*, 1991 *Nucl. Acids Res.* 19:3979-3986), which is hereby incorporated by reference), in certain embodiments without intronic sequences. Those skilled in the art, however, will appreciate that any of a number of other known promoters, such as the strong immunoglobulin, or other eukaryotic gene promoters

may also be used, including the EF1 alpha promoter, the murine CMV promoter, Rous sarcoma virus (RSV) promoter, SV40 early/late promoters and the beta-actin promoter.

The promoter may comprise a regulatable sequence such as the Tet operator sequence. This is extremely useful, for example, in cases where the gene products are affecting a result other than that desired and repression is sought.

Transcription termination sequences can also be utilized within the gene expression cassettes. Preferred termination sequences are, for instance, the bovine growth hormone terminator/polyadenylation signal (bGHpA) and the short synthetic polyA signal (SPA) of 50 nucleotides in length, defined as follows:

10 AATAAAAGATCTTTATTTTCATTAGATCTGTGTGTGTGTTTTTTTGTGTG (SEQ ID NO:4).

Further embodiments incorporate a leader or signal peptide into the transgene. A preferred leader is that from the tissue-specific plasminogen activator protein, tPA.

The following non-limiting Examples are presented to better illustrate the

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#### EXAMPLE 1

#### Construction and Rescue

An E1- Ad35-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad35 vector (a representative group B serotype) could be propagated in a group C E1-complementing cell line. The general strategy used to recover Ad35 as a bacterial plasmid is illustrated in Figure 3. Cotransformation of BJ5183 bacteria with purified wild-type Ad35 viral DNA and a second DNA fragment termed the Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad5 457 to 3402 with a unique Swa I site located in the deletion. The Ad35 sequences in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1. Pre-Adenovirus plasmid pAd35ΔE1 contains Ad35 sequences from 4 to 456 and bp 3403 to 34793.

To determine if pre-adenovirus plasmid pAd35ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a T-25 flask of PER.C6 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was never observed. Cells and media from the transfection were harvested at 14 days post transfection, freeze-thawed three times, clarified by centrifugation and used to infect new PER.C6 cells but no virus was ever amplified. Following multiple attempts, we have been unable to rescue and amplify pAd35ΔE1 in PER.C6 cells.

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#### **EXAMPLE 2**

## Insertion of Ad5 Orf 6 and Ad5 E4 into the Ad5 Genome

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in a Ad5/group C complementing cell line four additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. In the third strategy, the entire alternative serotype E4 coding region (not including the E4 promoter) was deleted and replaced with the Ad5 E4 coding region (not including the Ad5 E4 promoter) and, in the final strategy, the entire alternative serotype E4 coding and promoter region was deleted and replaced with the Ad5 E4 promoter and coding region. The configuration of the E4 regions generated by the four strategies is diagramed in Figure 4. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with purified wild-type viral DNA and the appropriately constructed ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of each pre-Ad plasmid, based on Ad35, is outlined below:

To construct pAd35 $\Delta$ E1 $\Delta$ E4Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a

bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31912 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 5). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5Orf6 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31913 and bp 34419 to bp 34793 with Ad5Orf6 cloned between bp 31913 and bp 34419.

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To construct pAd35ΔE1ΔOrf6Ad5Orf6 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 32081 and bp 32990 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-10. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 32081 and 32990 generating pNEBAd35-10Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and a deletion of E4 Orf6 from Ad35 bp 32082 to 32989 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct, Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 32081 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔOrf6Ad5Orf6. Pre-Adenovirus plasmid pAd35ΔE1ΔOrf6Ad5Orf6 contains Ad35

sequences from bp 4 to 456; bp 3403 to bp 32081 and bp 32990 to bp 34793 with Ad5Orf6 cloned between bp 32081 and bp 32990.

To construct pAd35ΔE1ΔE4Ad5E4 (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 E4), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-7. Next the Ad5 E4 coding region was generated by PCR and cloned between Ad35 bp 31838 and 34419 generating pNEBAd35-7Ad5E4-2 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34418 into which the Ad5 E4 coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5E4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5E4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34419 to bp 34793 with the Ad5 E4 coding region (Ad 5 bp 32914 to bp 35523) cloned between bp 31838 and bp 34419.

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To construct pAd35\(\text{E1}\text{DE4}\text{Ad5}\text{PE4}\) (An Ad35 pre-Ad plasmid containing an E1 deletion and a deletion of E4 coding region and promoter substituted with Ad5 E4 coding region and promoter), an Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31838 and bp 34660 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-8. Next the Ad5 E4 promoter and coding region was generated by PCR and cloned between Ad35 bp 31838 and 34660 generating pNEBAd35-8Ad5E4PC (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication,

ampicillin resistance gene, and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31839 to 34659 into which the Ad5 E4 promoter and coding region was introduced in an E4 parallel orientation. In this construct, the Ad5 E4 region is expressed using the Ad5 E4 promoter. The Ad35 sequences (bp 31599 to 31838 and bp 3403 to 3886) in the ITR cassette provide regions of homology with the purified Ad35 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the recombinant Ad35 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd35ΔE1ΔE4Ad5PE4. Pre-Adenovirus plasmid pAd35ΔE1ΔE4Ad5PE4 contains Ad35 sequences from bp 4 to 456; bp 3403 to bp 31838 and bp 34660 to bp 34793 with the Ad5 E4 promoter and coding region (Ad 5 bp 32914 to bp 35826) cloned between bp 31838 and bp 34660.

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# EXAMPLE 3 Rescue of pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4 into Virus

In order to determine if pre-adenovirus plasmids pAd35ΔE1ΔE4Ad5Orf6, pAd35 $\Delta$ E1 $\Delta$ Orf6Ad5Orf6, pAd35 $\Delta$ E1 $\Delta$ E4Ad5E4 and pAd35 $\Delta$ E1 $\Delta$ E4Ad5PE4 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmids were each digested with Pme I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for all construct. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect aT-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then

digested with *HindIII* and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/HindIII* prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### **EXAMPLE 4**

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Insertion of an Expression Cassette into pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4

In order to introduce a gag or SEAP expression cassette into the E1 region of the various Ad35 pre-Adenovirus plasmids described above (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4 and pAd35ΔE1ΔE4Ad5PE4) bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HTV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence (Figure 6), was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique SwaI site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being cloned into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad35 pre-Ad plasmids (pAd35ΔE1ΔE4Ad5Orf6, pAd35ΔE1ΔOrf6Ad5Orf6, pAd35ΔE1ΔE4Ad5E4, pAd35ΔE1ΔE4Ad5PE4), linearized in the E1 region by digestion with Swa I, resulted in the generation of corresponding Ad35 gag-containing pre-Adenovirus plasmids (pAd35ΔE1gagΔE4Ad5Orf6, pAd35ΔE1gagΔOrf6Ad5Orf6, pAd35ΔE1gagΔE4Ad5E4, and pAd35ΔE1gagΔE4Ad5PE4) by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad35 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence

[Figure 7] was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. The transgene was then recombined into the various Ad35 backbones generating pAd35ΔE1SEAPΔE4Ad5Orf6, pAd35ΔE1SEAPΔE4Ad5E4, and pAd35ΔE1SEAPΔE4Ad5PE4 as described above for the gag transgene.

All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

## 15 EXAMPLE 5

In vivo Transgene Expression

#### A. Immunization

Female mice were between 4-10 weeks old. The total dose of each vaccine was
suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each animals with a volume of 50 μL per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, Franklin Lakes, NJ). The rhesus macaques and African green monkeys were between 2-5 kg in weight. For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National
Research Council.

#### B. SEAP Assay

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Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5  $\mu$ L aliquots of each serum were mixed with 45  $\mu$ L of kit-supplied dilution buffer in a 96-well white DYNEX plate.

Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey or mouse serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

#### C. Rodent Results

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In the first mouse experiment, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (2) 10^10 vp Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1SEAP. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 8. Results indicate that (1) the Ad35 constructs are all capable of expressing the SEAP transgene and that (2) the introduction of Ad5Orf6 sequence where the deleted Ad35E4 was did not significantly affect the transgene expression relative to Ad35ΔE1SEAP. Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 also yielded a similar expression profile as Ad35ΔE1SEAP. The levels of SEAP in the serum dropped after day 2 and were at background levels by day 12.

The second mouse experiment evaluates the effect of a full Ad5E4 replacement instead of an Ad5Orf6 substitution for the Ad35 E4 cassette. Here, cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4; or (5) 10^10 vp Ad35ΔE1SEAPΔE4Ad5PE4. The introduction of Ad5E4 or Ad5PE4 resulted in comparable if not, slightly improved expression levels compared to the vector with the Ad5Orf6 insertion (Figure 9). The peak levels for the Ad35 constructs are lower than those produced by Ad5SEAP (at least 10-fold).

#### D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; or (3) 10^11 vp Ad35\Delta E1SEAP\Delta E4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results for the individual monkeys are shown in Figures 10A-B. Results indicate that the peak level of SEAP product produced by the alternative adenovirus serotype was lower than but were within 3-fold of that of MRKAd5SEAP at the same

high dose level of 10^11 vp. The levels observed from the Ad35 vector were about 50-fold higher than those observed using 10^9 vp of MRKAd5SEAP. The levels of SEAP in the serum dropped after day 10 and were close to background as early as day 15.

A separate experiment using African green monkeys was conducted to examine the effect of the additional E3 deletion or the full Ad5E4 substitution on in vivo gene expression. In here, cohorts of 2-3 African green macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^10 vp MRKAd5-SEAP; (3) 10^9 vp MRKAd5-SEAP; (4) 10^10 vp Ad35ΔE1SEAPΔE4Ad5Orf6; (5) 10^10 vp Ad35ΔE1SEAPΔE4Ad5E4. Results (Figure 11) indicate that the peak levels of SEAP product produced by Ad35ΔE1SEAPΔE3ΔE4Ad5Orf6 and Ad35ΔE1SEAPΔE4Ad5E4 were comparable if not, slightly improved compared to Ad35ΔE1SEAPΔE4Ad5Orf6.

#### EXAMPLE 6

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#### 15 In vivo Immunogenicity

#### A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; or (2) 10^11 vp of Ad35\Delta 1 gag\Delta E4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson). Sera and peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### 30 B. ELISPOT Assay

The IFN-γ ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen *et al.*, 2001 *J. Virol.* 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50 μL of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs)

were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50 μL of media or the gag peptide pool at 8 μg/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10<sup>6</sup> cell input.

## C. Intracellular Cytokine Staining

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To 1 ml of 2 x 10<sup>6</sup> PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1  $\mu$ g/mL. For gag-specific stimulation, 10  $\mu$ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20  $\mu L$  of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson, Franklin Lakes, NJ); and 20  $\mu$ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu L$ 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4+ and CD8+ populations, and for both mock and gag-peptide reaction tubes of a sample.

#### 30 D. Results

PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay. Results (Table 1) indicate that the Ad35ΔE1gagΔE4Ad5Orf6 is able to induce in non-human primates significant levels of gag-specific T cells. After a single dose (wk 4), the Ad35-induced responses were about 5-fold lower than that of MRKAd5-HIV1 gag. After the second dose (wk

8), the responses between both cohorts were comparable; the differences became pronounced in the succeeding time points.

Table 1. Gag-specific T cell response in monkeys immunized with MRKAd5-HIV1gag and Ad35ΔE1gagΔE4Ad5Orf6. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mock) or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

Gгр	Vaccine Wk 0, Wk 4	Monkey ID	Pre		Wk 4		Wk 8		Wk 12		Wk 16	
			Mock	Gag H		Gag H	Mock	Gag H	Mock	Gag H	Mock	Gag
1	MRKAd5-HIV1 gag 10^11 vp	00C018 00C034 00C058	1 0 4	5 4 4	13 5 3	1025 219 1086	0 5 0	824 404 440	3 0 0	.753 491 439	1 1 0	533 350 599
2	Ad35AE1gagAE4Ad5Ori6 10^11 vp	00D045 00D067 00D068 00D054 00D075 00D073	1 1 3 3 14	1 4 4 15 5 26	3 5 10 10 18 1	168 89 34 195 275 241	5 0 5 0 13 3	645 103 365 501 716 485	4 0 3 3 3 3	178 76 143 350 158 278	0 0 0	91 19 95 124 103 148
	Naïve	00D087	1	1	3	3	8	54	3	5	3	<u> 1</u>

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Intracellular IFN-γ staining analyses of PBMC collected at wk 8 suggest that the Ad35-based vaccine is able to induce both HIV-specific CD4+ and CD8+ T cells (Table 2).

Table 2. Characterization of the gag-specific T cells in monkeys immunized with MRKAd5-HIV1gag and Ad35ΔE1gagΔE4Ad5Orf6. Shown are the percentages of CD3+ T cells that are either gag-specific CD4+ or gag-specific CD8+ cells. These values were corrected for mock values (<0.02%).

Grp	Vaccine	Monkey	Wk 8			
Gip	Wk 0, Wk 4	ID	%CD4+CD3+	%CD8+CD3+		
1	MRKAd5-HIV1 gag 10^11 vp	00C018 00C034 00C058	0.08 0.09 0.03	0.37 0.06 0.21		
2	Ad35∆E1gag∆E4Ad5Orf6 10^11 vp	00D045 00D067 00D068 00D054 00D075 00D073	0.06 0.02 0.15 0.05 0.08 0.09	0.08 0.02 0.02 0.08 0.05 0.06		

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In a separate experiment, 3 different Ad35 constructs expressing HIV-1 gag were evaluated for their immunogenicity in macaques. Here, cohorts of 3 macaques were given immunizations at wk 0 and 4 of either of the following vectors: (1) 10^10 vp Ad35ΔE1gagΔE4Ad5Orf6; (2) 10^10

vp Ad35ΔE1gagΔE3ΔE4Ad5Orf6; or (3) 10^10 vp Ad35ΔE1gagΔE4Ad5E4. The levels of T cell immunity induced by all 3 vectors were comparable at this stage (Table 2), suggesting that the additional E3 deletion or full Ad5E4 substitution does not appear to impair the immunogenic properties of the vector.

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Table 3. Gag-specific T cell response in monkeys immunized with several Ad35ΔE1ΔE4-based vectors. Shown is the number of spot-forming cells per million PBMC following incubation in the absence (mocK0 or presence of Gag H peptide pool. The H pool consisted of 20-aa peptide overlapping by 10 aa and encompassing the entire gag sequence.

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Grp	Vaccine	Monkey	Pre		Wk 4		Wk 8	
GIP	Wk 0, Wk 4	ID	Mock	Gag H	Mock	Gag H	Mock	Gag H
<del></del>	Ad35∆E1gag∆E4Ad5Orf6	00C047	4	1	0	20	0	189
'	10^10vp	00C157	8	5	1	81	1	833
	10 104β	00C078	3	1	0	46	4	349
2	Ad35∆E1gag∆E3∆E4Ad5Orf8	000091	1	1	1	118	3	315
2	10^10vp	00C122	3	loi	0	31	1	138
	ιστιούρ	00D177	3	3	1	45	1	64
3	Ad35∆E1gag∆E4Ad5E4	00D018	3	19	29	120	23	193
	10^10vp	00D046	8	5	1	21	10	143
	10 1049	00D063	3	4	0	63	4	371
Naîve	none	00D363	0	5	ND	ND	0	0

## EXAMPLE 7 Construction and Rescue of pAd24ΔE1.

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An E1- Ad24-based pre-adenovirus plasmid was constructed in order to determine whether an E1- Ad24 vector (a representative group D serotype) could be propagated in an Ad5/group C E1-complementing cell line. Since at the time the vector construction was initiated the complete sequence of Ad24 (see Figures 16A-1 through 16A-10; subject of copending application serial no. 60/455, 312, filed March 17, 2003) was unknown we took advantage of some sequence homology between Ad24 and Ad17. The general strategy used to recover Ad24 as a bacterial plasmid is illustrated in Figure 12 and described below. Cotransformation of BJ5183 bacteria with purified wild-type Ad24 viral DNA and a second DNA fragment termed the Ad17 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome (Accession No. AF108105) separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17

(bp 415 to 3372) with a unique Swa I site located in the deletion. The Ad17 sequences in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which recombination can occur. The ITR cassette was also designed to contain unique restriction enzyme sites (Pme I) located at the end of the viral ITR's so that digestion will release the Ad24 genome from plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd24ΔE1. pAd24ΔE1 contains Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24). PAd24ΔE1 contains the coding sequences for all Ad24 virion structural proteins that constitute its serotype specificity. This approach can be used to circularize any group D serotype into plasmid form which has sufficient homology to Ad17.

To determine if pre-adenovirus plasmid pAd24ΔE1 could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. *Pme* I digestion releases the viral genome from the plasmid sequences allowing viral replication to occur after entry into 293 cells. Viral cytopathic effect (CPE), indicating that virus replication and amplification is occurring, was very slow to arise. Following multiple attempts, we were successful at rescuing and amplifying Ad24ΔE1 but the virus grew to lower titers and took more passages to amplify than a similar Ad5 based vector. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *Hind*III and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with *Pme1/Hind*III prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

#### **EXAMPLE 8**

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## Insertion of Ad5 Orf 6 into the E1 region of Ad24

In order to determine if the insertion of Ad5 E4 Orf6 into the Ad24 genome would allow more efficient propagation in a group C E1 complementing cell line we constructed an Ad24 based pre-adenovirus plasmid containing Ad5 Orf6 in the E1 region. In order to introduce Ad5 Orf6 in to the E1 region of pAd24ΔE1, bacterial recombination was used. An Ad5 Orf6 transgene consisting of the Ad5 Orf6 coding region flanked by the HCMV promoter and pA was cloned into the E1 deletion in an Ad17 shuttle vector (a precursor to the Ad17 ITR cassette). The Ad5 Orf6 transgene was cloned between bp 414 and 3373 in the E1 anti-parallel

orientation. The shuttle vector containing the Ad5 Orf6 transgene was digested to generate a DNA fragment consisting of the transgene flanked by Ad17 sequences (bp 4 to 414 and bp 3373 to 4580) and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd24ΔE1, which had been linearized in the E1 region by digestion with *Swa*I, resulted in the generation of pAd24ΔE1Ad5Orf6 by homologous recombination (Figure 13). Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1Ad5Orf6.

In order to determine if pre-adenovirus plasmid pAd24ΔE1Ad5Orf6 could be rescued into virus and propagated in an Ad5/group C E1 complementing cell line, pAd24ΔE1Ad5Orf6 was digested with Pme I and transfected into a 6 cm dish of 293 cells using the calcium phosphate co-precipitation technique. PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after entry into 293 cells. Once complete viral cytopathic effect (CPE) was observed at approximately 7-10 days post transfection, the infected cells and media were freeze/thawed three times and the cell debris pelleted. The virus was amplified in two additional passages in 293 cells and then purified from the final infection by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the virus, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The endlabeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products from the pre-plasmid (that had been digested with Pme1/HindIII prior to labeling). The expected sizes were observed, indicating that the virus had been successfully rescued.

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#### **EXAMPLE 9**

## Insertion of Ad5 Orf 6 into the E4 region of Ad24

To refine the strategy of including Ad5 Orf6 in the genome of an alternative serotype so that propagation could take place in an Ad5/group C complementing cell line two additional strategies were developed. In the first strategy, the entire alternative serotype E4 region (not including the E4 promoter) was deleted and replaced with Ad5 Orf6. In the second strategy, just the alternative serotype Orf6 gene was deleted and replaced with Ad5 Orf6. The configuration of the E4 regions generated by the two strategies is diagramed in Figure 14. For each of these strategies the desired pre-Adenovirus plasmid was generated by bacterial recombination. Cotransformation of BJ 5183 bacteria with pAd24ΔOrf6BstZ17I and the

appropriately constructed Ad24 E4 shuttle plasmid resulted in the generation of the desired Ad24 based pre-Ad plasmid. PAd24ΔOrf6BstZ17I, a derivative of pAd24ΔE1, was constructed so that the E4 region in the Ad24 pre-Ad plasmid could be easily modified using bacterial recombination. PAd24ΔOrf6BstZ17I contains a deletion in the E4 region from Ad24 bp 32373 to bp 33328 with a unique BstZ17I site located at the position of the deletion. The complete sequence of pAd24ΔOrf6BstZ17I consists of Ad17 sequences from bp 4 to 414 and from bp 3373 to 4580, Ad24 bp 4588 to 32372 and from 33329 to 34529, and Ad17 bp 34469 to 35098 (bp numbers refer to the wt sequence for both Ad17 and Ad24).

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To construct pAd24ΔE1ΔE4Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed by digesting pAd24ΔE1 with *PmeI* and *BsrGI* and cloning the restriction fragment representing the E4 region (bp 31559 to bp 35164) into pNEB193, generating pNEBAd24E4. PNEBAd24E4 was then digested with *AccI* and *Eco*NI to remove the E4 coding sequences and ligated with an oligo designed to contain *BglII* and *XhoI* sites (underlined) (5'

ACTCGAGATGTATAGATCT (SEQ ID NO: 6); 5' CTAGATCTATACATCTCGAG (SEQ ID NO: 7)), generating pNEBAd24ΔE4. PNEBAd24ΔE4 was then digested with BgIII and XhoI and ligated with the Ad5 Orf6 gene, which was PCR amplified, generating pNEBAd24ΔE4Ad5Orf6. The PCR primers used to amplify the Ad5 Orf6 gene (5' GCACAGATCTTTGCTTCAGGAATATG (SEQ ID NO: 8); 5'

GAGAACTCGAGGCCTACATGGGGGTAGAG (SEQ ID NO: 9)) were designed to contain BglII and XhoI sites (underlined above) for ligation with the pNEBAd24DE4 fragment. In the final step pNEBAd24ΔE4Ad5Orf6 E4 shuttle plasmid was digested with PvuI and PmeI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with E4 shuttle fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔE4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as pre-adenovirus plasmid pAd24ΔE1ΔE4Ad5Orf6.

To construct pAd24ΔE1ΔOrf6Ad5Orf6 (An Ad24 pre-Ad plasmid containing an E1 deletion and a deletion of E4 Orf6 substituted with Ad5 Orf6), an Ad24 E4 shuttle plasmid was constructed in which the Ad24 Orf6 gene was replaced by Ad5 Orf6. To do this the *Eco*R1 restriction fragment representing bp 32126 to bp 33442 of the Ad24 genome (encompassing the E4 Orf6 coding region), was subcloned into the *Eco*RI site in pNEB193, generating pNEBAd24Orf6. In order to delete the E4 Orf6 gene in pNEBAd24Orf6 and replace it with Ad5 Orf6, pNEBAd24Orf6 was digested with *Sty*I and treated with Klenow to blunt the ends and then

digested with to EagI. The desired pNEBAd24Orf6 fragment was then ligated with a PCR product representing the Ad5 Orf6 gene from Ad5 bp 33193 to bp 24125, generating pNEBAd24ΔOrf6Ad5Orf6. The PCR primers used to generate the Ad5 Orf6 fragment (5'CGAGACGCCGACGCAGATCTGTTTG (SEQ ID NO: 10);

5 'GAAGTCCCGGGCTACATGGGGGTAG (SEQ ID NO: 11)) were designed to contain EagI and SmaI sites (underlined above) for ligation with the pNEBAd24Orf6 fragment. In the final step pNEBAd24ΔOrf6Ad5Orf6 was digested with EcoRI, the restriction fragments were size fractionated by agarose gel electrophoresis and the desired fragment containing Ad5Orf6 flanked by Ad24 sequences was gel purified. Cotransformation of BJ 5183 bacteria with the EcoRI fragment and pAd24ΔOrf6BstZ17I, which had been linearized in the E4 region by digestion with BstZ17I, resulted in the generation of pAd24ΔE1ΔOrf6Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis and one clone was selected as preadenovirus plasmid pAd24ΔE1ΔOrf6Ad5Orf6.

#### EXAMPLE 10

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## Rescue of pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, into Virus

In order to determine if pre-adenovirus plasmids pAd24\Delta E1\Delta E4Ad5Orf6, pAd24AE1AOrf6Ad5Orf6, could be rescued into virus and propagated in a group CE1 complementing cell line, the plasmids were each digested with Pme I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique; (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc.). PmeI digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring, was observed for both constructs. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Approximately 1 ml of the cell lysate was used to infect T-225 flasks of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with HindIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared

with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with Pme1/HindIII prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### **EXAMPLE 11** 5

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## Comparison of the Growth Kinetics of Ad24 based vectors.

In order to compare the growth kinetic of Ad24ΔE1, Ad24ΔE1Ad5Orf6, Ad24ΔE1ΔE4Ad5Orf6 and Ad24ΔE1ΔOrf6Ad5Orf6 one step growth curves were preformed (Figure 15). PER.C6 cells in 60 mm dishes were infected at 1 vp per cell with either Ad24 $\Delta$ E1, Ad24ΔE1Ad5Orf6, Ad24ΔE1ΔE4Ad5Orf6 or Ad24ΔE1ΔOrf6Ad5Orf6. Cells and media were then harvested at various times post infection, freeze thawed three times and clarified by centrifugation. The amount of virus present in the samples was determined by quantitative PCR and is illustrated in Figure 15. This study demonstrates that Ad24 vectors that incorporate Ad5 Orf6 have a distinct growth advantage over Ad24ΔE1 in PER.C6 cells. The instant invention can be practiced with recombinant Ad24 vectors absent a heterologous Orf 6 region where the E1-complementing cell line expresses an Ad24 E1 region or, alternatively, E1 and E4 regions of the same serotype (such as Ad5E1/E4-expressing cell lines).

#### **EXAMPLE 12**

Insertion of an Expression Cassette into pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6, In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, respectively) into the E1 region of the Ad24 pre-Adenovirus plasmids described above (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6) bacterial recombination was used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVgagBGHpA. The ITR cassette contains sequences from the right (bp 34469 to 35098) and left (bp 4 to 414 and bp 3373 to 4580) end of the Ad17 genome separated by plasmid sequences containing a bacterial origin of replication and an Ampicillin 30 resistance gene. The ITR cassette contains a deletion of E1 sequences from Ad17 (bp 415 to 3372) with a unique Swa I site located in the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the Swal site in pABSAd17-3. This cloning step resulted in the gag expression cassette being 35

cloned into the E1 deletion between bp 414 and 3373 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 gag-containing pre-Adenovirus plasmids (pAd24ΔE1gagΔE4Ad5Orf6, pAd24ΔE1gagΔOrf6Ad5Orf6) by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad24 pre-Ad plasmids containing a SEAP 10 expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was cloned into the E1 deletion in Ad17 shuttle plasmid, pABSAd17-3, generating pABSAd17HCMVSEAPBGH. The SEAP expression cassette was obtained from a previously 15 constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pABSAd17-3. The shuttle vector containing the SEAP transgene was digested to generate a DNA fragment consisting of the SEAP expression cassette flanked by Ad17 bp 4 to 414 and bp 3373 to 4580 and the fragment was purified after electrophoresis on an agarose gel. 20 Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and one of the Ad24 pre-Ad plasmids (pAd24ΔE1ΔE4Ad5Orf6, pAd24ΔE1ΔOrf6Ad5Orf6,), linearized in the E1 region by digestion with Swa I, resulted in the generation of the corresponding Ad24 SEAP-containing pre-Adenovirus plasmids (pAd24ΔE1SEAPΔE4Ad5Orf6, pAd24ΔE1SEAPΔOrf6Ad5Orf6) by

homologous recombination. Potential clones were screened by restriction analysis. All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### **EXAMPLE 13**

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30 In Vivo Immunogenicity

#### A. Immunization

Cohorts of 3-6 animals were given intramuscular injections at wk 0 and wk 4 of either of the following constructs: (1) 10^11 vp MRKAd5-HIV1 gag; (2) 10^10 vp MRKAd5-HIV1 gag; (3) 10^11 vp of Ad24ΔE1gagΔOrf6Ad5Orf6; (4) 10^10 vp of

Ad24ΔE1gagΔOrf6Ad5Orf6; or (5) 10^10 vp of Ad24ΔE1gagΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points (typically 4 wk intervals) during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

#### B. ELISPOT Assay

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The IFN- $\gamma$  ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen et al., 2001 *J. Virol.* 75(2):738-749; Casimiro et al., 2002 *J. Virol.* 76:185-94), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50  $\mu$ L of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower size cut-off set at 80 femtoliters ("fL"). Either 50  $\mu$ L of media or the gag peptide pool at 8  $\mu$ g/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to 10<sup>6</sup> cell input.

## 25 C. Intracellular Cytokine Staining

To 1 ml of 2 x 10<sup>6</sup> PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1 μg/mL. For gag-specific stimulation, 10 μL of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 μL per tube anti-hCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson);

and 20  $\mu$ L anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750  $\mu$ L 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1  $\mu$ g of FITC-anti-hIFN- $\gamma$ , clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4<sup>+</sup> and CD8<sup>+</sup> populations, and for both mock and gag-peptide reaction tubes of a sample.

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#### D. Anti-p24 ELISA

A modified competitive anti-p24 assay was developed using reagents from the Coulter p24 Antigen Assay kit (Beckman Coulter, Fullerton, CA). Briefly, to a 250-μL serum sample, 20 μL of Lyse Buffer and 15 μL of p24 antigen (9.375 pg) from the Coulter kit were added. After mixing, 200 μL of each sample were added to wells coated with a mouse anti-p24 mAb from the Coulter kit and incubated for 1.5 hr at 37°C. The wells were then washed and 200 μL of Biotin Reagent (polyclonal anti-p24-biotin) from the Coulter kit was added to each well. After a 1 hr, 37°C incubation, detection was achieved using strepavidin-conjugated horseradish peroxidase and TMB substrate as described in the Coulter Kit. OD450nm values were recorded. A 7-point standard curve was generated using a serial 2-fold dilution of serum from an HIV-seropositive individual. The lower cut-off for the assay is arbitrarily set at 10 milli Merck units/mL (mMU/mL) defined by a dilution of the seropositive human serum. This cutoff falls at approximately 65% of the maximum bound control signal which corresponds to that obtained with the diluent control only and with no positive analyte.

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#### E. Results

PBMCs collected at regular 4-wk intervals were analyzed in an ELISPOT assay (Figure 17). Both Ad24ΔE1gagΔOrf6Ad5Orf6 and Ad24ΔE1gagΔE4Ad5Orf6 were able to induce significant levels of gag-specific T cells in non-human primates. At 10^11 vp dose level, the Ad24-induced responses were within 2-3-fold of those of MRKAd5-HIV1 gag. Both Ad24 vectors were also able to induce detectable levels of gag-specific T cells at 10^10 vp but were lower than those observed using MRKad5gag at the same dose.

PBMCs collected at wk 12 from the vaccinees were analyzed for intracellular IFN-γ staining after the priming immunizations. The assay results provided information on the relative amounts of CD4<sup>+</sup> and CD8<sup>+</sup> gag-specific T cells in the peripheral blood (Figure 18). The

results indicated that the prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cells.

## F. Humoral Immune Responses

The Ad24-based vaccine vector was able to generate detectable levels of circulating anti-gag antibodies at the reasonably high dose level (Figure 19). No detectable titers were observed at equal to or lower than 10^10 vp, suggesting the existence of a dose-dependent response.

#### EXAMPLE 14

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In Vivo Transgene Expression

#### A. Immunization

Cohorts of 5 C3H/HeN mice were given single intramuscular injections of one of the following vectors: (1) 10^10 vp Ad24ΔE1SEAPΔE4Ad5Orf6; (2) 10^10 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; (3) 10^10 vp MRKAd5SEAP; and (4) 10^9 vp MRKAd5SEAP. Female mice were between 4-10 weeks old. The total dose of each vaccine was suspended in 0.1 mL of buffer. The vectors were given to both quadriceps of each of the animals with a volume of 50 uL per quad and using 0.3-mL 28G1/2 insulin syringes (Becton-Dickinson, Franklin Lakes, NJ). For the primates, the total dose of each vaccine was suspended in 1 mL of buffer. The monkeys were anesthetized (ketamine/xylazine mixture) and the vaccines were delivered i.m. in 0.5-mL aliquots into two muscle sites using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Serum samples were collected at defined intervals and stored frozen until the assay date. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### B. SEAP Assay

Serum samples were analyzed for circulating SEAP levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 uL aliquots of each serum were mixed with 45 uL of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the wells for 30 minutes at 65 °C.

Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorder using DYNEX luminometer. RLU readings are converted to ng/mL SEAP using a log-log regression analyses.

#### 5 C. Rodent Results

Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 20. Results indicate that (1) both Ad24 constructs are all capable of expressing the SEAP transgene in vivo to comparable levels; and that (2) the level of expression achieved using the Ad24 vectors are comparable to that of Ad5 at 10-fold lower dose. The levels of SEAP in the serum dropped dramatically after day 2 and were at background levels by day 12.

#### D. Primate Results

Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the following vectors: (1) 10^11 vp MRKAd5-SEAP; (2) 10^9 vp MRKAd5-SEAP; (3) 10^11 vp Ad24ΔE1SEAPΔOrf6Ad5Orf6; or (4) 10^11 vp Ad24ΔE1SEAPΔE4Ad5Orf6. Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 21.

Results indicate that the peak levels of SEAP product produced by adenovirus serotype 24 were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 21). The levels observed with adenovirus serotype 24 are generally 50-fold higher than those observed using 10^9 vp of MRKAd5. The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that adenovirus serotype 24 is very efficient in expressing a transgene following intramuscular administration in a primate.

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#### EXAMPLE 15

# Construction of pMRKAd24\Delta E1\Delta E4Ad5Orf6

To construct pMRKAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6 (An Ad24 pre-Ad plasmid, composed entirely of Ad24 sequence and containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), an Ad24 ITR cassette was constructed containing sequences from the right (bp 31978 to 32264 and bp 34713 to 35164) and left (bp 4 to 450 and bp 3364 to 3799) end of the Ad24 genome separated by plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd24-4. Next the Ad5 Orf6 open reading frame (Ad5 bp 31192 to bp 34078) was generated by PCR and cloned between Ad24 bp 32264 and 34713 generating

pNEBAd24E-Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad24 bp 451 to 3363 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad24 bp 32265 to 34712 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5 Orf6 expression is driven by the Ad24 E4 promoter. The Ad24 sequences (bp 31978 to 32264 and bp 3464 to 3799) in the ITR cassette provide regions of homology with the purified Ad24 viral DNA in which bacterial recombination can occur following cotransformation into BJ 5183 bacteria (Figure 22). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion will release the recombinant Ad24 genome from plasmid sequences. Potential clones will be screened by restriction analysis and one clone was selected as pMRKAd24ΔE1ΔE4Ad5Orf6. Pre-Adenovirus plasmid pMRKAd24\Delta E1\Delta E4Ad5Orf6 should contain Ad24 sequences from bp 4 to 450; bp 3364 to bp 32264 and bp 34713 to bp 35164 with Ad5Orf6 cloned between bp 32264 and bp 34713. The bp numbering in the above description refers to the wt sequence for both Ad24 and Ad5.

#### EXAMPLE 16

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# 20 Insertion of HIV-1 gag and SEAP transgenes into pAd24ΔE1ΔE4Ad5Orf6

In order to introduce a gag or SEAP expression cassettes into the E1 region of pMRKAd24ΔE1ΔE4Ad5Orf6, bacterial recombination will be used. An HIV-1 gag expression cassette will consist of the following: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, in the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2 (a precursor to the Ad24 ITR cassette described above), generating pNEBAd24CMVgagBGHpA. PNEBAd24-2 contains Ad24 sequences from the left end of the genome (bp 4 to 450 and bp 3364 to 3799) that define the E1 deletion. The gag expression cassette will be obtained from a previously constructed plasmid and cloned into the E1 deletion between bp 450 and 3364 in the E1 parallel orientation. The shuttle vector containing the gag transgene will be digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad24 bp 4 to 450 and bp 3364 to 3799 and the fragment will be purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pMRKAd24ΔE1ΔE4Ad5Orf6 which was linearized in the E1 region by digestion with SwaI, should result in the generation of Ad24 gag-

containing pre-Adenovirus plasmids pMRKAd24ΔE1gagΔE4Ad5Orf6 by homologous recombination. Potential clones will be screened by restriction analysis.

A similar strategy will be used to generate Ad24 pre-Ad plasmids containing a SEAP expression cassette. In this case, a SEAP expression cassette will consist of: 1) the immediate early gene promoter from the human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence cloned into the E1 deletion of an Ad24 shuttle plasmid, pNEBAd24-2, generating pNEBAd24CMVSEAPBGHpA. The transgene will then be recombined into pMRKAd24 $\Delta$ E1 $\Delta$ E4Ad5Orf6 as described above for the gag transgene.

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# EXAMPLE 17 In Vivo Immunogenicity

#### A. Immunization

Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the *Guide for Care and Use of Laboratory Animals*, Institute of Laboratory Animal Resources, National Research Council.

#### 25 B. T Cell Responses

Ad24 Vaccine Vector as a Heterologous Booster: Cohort of 4 rhesus macaques was immunized initially with 3 doses (wk 0, 4, 26) of either 10<sup>7</sup> or 10<sup>9</sup> vp of MRKAd5-gag (see, PCT/US01/28861, published March 21, 2002) or MRKAd6-gag. At wk 56, the animals received a booster vaccine of 10<sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6. A separate cohort of naïve animals received a single dose of the booster vaccine. The results of the IFN-γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 23. It is apparent that the Ad24 HIV vectors can be utilized to amplify the existing pools of HIV-specific T cells. The increases in the levels of gag-specific T cells from the pre-boost levels to those measured at 4 wks post boost were consistently larger than the levels induced by the same booster vaccine in naïve animals. PBMCs from the vaccinees of the heterologous MRKAd5/MRKAd6-Ad24 boost

regimen were analyzed for intracellular IFN-γ staining after the priming immunizations (wk 60). The assay results provided information on the relative amounts of CD4<sup>+</sup> and CD8<sup>+</sup> gag-specific T cells in the peripheral blood (Figure 24). The results indicated that heterologous prime-boost immunization approach was able to elicit in rhesus macaques both HIV-specific CD4+ and CD8+ T cells.

Ad24 Vaccine Vector as a Heterologous Primer: In a separate study, a cohort of 3 rhesus macaques was immunized initially with 2 doses (wk 0, 4) of 10<sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6 and boosted at wk 24 with 10<sup>7</sup> vp of MRKAd5-gag. The low dose of MRKAd5-gag is selected to mimic the effect of pre-existing neutralizing immunity to the vector in a subject. A separate cohort of naïve animals was given a single dose of 10<sup>7</sup> vp MRKAd5-gag. The results of the IFN-γ ELISPOT analyses of PBMC collected during the course of the studies are shown in Figure 25.

The Ad24-based vaccine was able to prime effectively for HIV-specific T cell responses in macaques. Boosting with a low dose MRKAd5-gag resulted in a significant increase in the levels of gag-specific T cells. The increases in 2 out of 3 animals exceed the levels typically observed after treatment of naïve animals with the same low dose of MRKAd5-gag.

#### **EXAMPLE 18**

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#### 20 Construction of pAd34ΔE1ΔE4Ad5Orf6

To generate an E1- Ad34 based vector that can propagate in existing group C/Ad5 E1 complementing cell lines (293, PER.C6), Ad5 Orf6 was inserted in place of the native E4 region. Since at the time, the complete sequence of Ad34 (see Figures 28A-1 to 28A-9; subject of copending application serial no. 60/458,825, filed March 28, 2003) was unknown, advantage was taken of the sequence homology between Ad34 and Ad35 in order to construct the Ad34 pre-Adenovirus plasmid. Cotransformation of BJ 5183 bacteria with purified wild-type Ad34 viral DNA and the appropriately constructed Ad35 ITR cassette resulted in the circularization of the viral genome by homologous recombination. The construction of the pre-Ad plasmid based on Ad34, is outlined below:

To construct pAd34ΔE1ΔE4Ad5Orf6 (An Ad34 pre-Ad plasmid containing an E1 deletion and an E4 deletion substituted with Ad5 Orf6), we utilized an Ad35 ITR cassette. We anticipated that sequence homology between Ad34 and Ad35 would allow homologous recombination to occur. The Ad35 ITR cassette was constructed containing sequences from the right (bp 31599 to 31913 and bp 34419 to 34793) and left (bp 4 to 456 and bp 3403 to 3886) end of the Ad35 genome (see Figures 2A-1 to 2A-10) separated by plasmid sequences containing a

bacterial origin of replication and an ampicillin resistance gene. The four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd35-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad35 bp 31913 and 34419 generating pNEBAd35-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad35 bp 457 to 3402 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad35 bp 31914 to 34418 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the Ad35 E4 promoter. The Ad35 sequences (bp 31599 to 31913 and bp 3403 to 3886) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 26). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

#### **EXAMPLE 19**

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#### 20 Rescue of pAd34ΔE1ΔE4Ad5Orf6 into Virus

In order to determine if pre-adenovirus plasmid pAd34AE1AE4Ad5Orf6, could be rescued into virus and propagated in a group C E1 complementing cell line, the plasmid was digested with *Pme* I and transfected into T-25 flasks of PER.C6 cells using the calcium phosphate co-precipitation technique (Cell Phect Transfection Kit, Amersham Pharmacia Biotech Inc). *Pme*I digestion releases the viral genome from plasmid sequences allowing viral replication to occur after cell entry. Viral cytopathic effect (CPE), indicating that virus replication and amplification was occurring was observed following transfection. When CPE was complete, approximately 7-10 days post transfection, the infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation.

Approximately 1 ml of the cell lysate was used to infect a T-225 flask of PER.C6 cells at 80-90% confluence. Once CPE was reached, infected cells and media were harvested, freeze/thawed three times and the cell debris pelleted by centrifugation. Clarified cell lysates were then used to infect 2-layer NUNC cell factories of PER.C6 cells. Following complete CPE, the virus was purified by ultracentrifugation on CsCl density gradients. In order to verify the genetic structure of the rescued viruses, viral DNA was extracted using pronase treatment

followed by phenol chloroform extraction and ethanol precipitation. Viral DNA was then digested with *Hin*dIII and treated with Klenow fragment to end-label the restriction fragments with P33-dATP. The end-labeled restriction fragments were then size-fractionated by gel electrophoresis and visualized by autoradiography. The digestion products were compared with the digestion products of the corresponding pre-Adenovirus plasmid (that had been digested with *Pme1/Hind*III prior to labeling) from which they were derived. The expected sizes were observed, indicating that the viruses had been successfully rescued.

#### EXAMPLE 20

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### Insertion of an Expression Cassette into pAd34ΔE1ΔE4Ad5Orf6

In order to introduce a gag or SEAP expression cassette (see Figures 6 and 7, respectively) into the E1 region of pAd34ΔE1ΔE4Ad5Orf6, bacterial recombination was again used. A gag expression cassette consisting of the following: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human immunodeficiency virus type 1 (HIV-1) gag (strain CAM-1; 1526 bp) gene, and 3) the bovine growth hormone polyadenylation signal sequence, was cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2 (a precursor to the Ad35 ITR cassettes described above), generating pNEBAd35CMVgagBGHpA. pNEBAd35-2 contains Ad35 sequences from the left end of the genome (bp 4 to 456 and bp 3403 to 3886) with a unique SwaI site between bp 456 and 3403 at the position of the deletion. The gag expression cassette was obtained from a previously constructed shuttle plasmid by EcoRI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the SwaI site in pNEBAd35-2. This cloning step resulted in the gag expression cassette being inserted into the E1 deletion between bp 456 and 3403 in the E1 parallel orientation. The shuttle vector containing the gag transgene was digested to generate a DNA fragment consisting of the gag expression cassette flanked by Ad35 bp 4 to 456 and bp 3403 to 3886 and the fragment was purified after electrophoresis on an agarose gel. Cotransformation of BJ 5183 bacteria with the shuttle vector fragment and pAd34\Delta E1\Delta E4Ad5Orf6, linearized in the E1 region by digestion with Swa I, resulted in the generation of the Ad34 gag-containing pre-Adenovirus plasmid pAd34ΔE1gagΔE4Ad5Orf6 by homologous recombination. Potential clones were screened by restriction analysis.

A similar strategy was used to generate Ad34 pre-Ad plasmids containing a SEAP expression cassette. In this case a SEAP expression cassette consisting of: 1) the immediate early gene promoter from human cytomegalovirus, 2) the coding sequence of the human placental SEAP gene, and 3) the bovine growth hormone polyadenylation signal sequence was

cloned into the E1 deletion in Ad35 shuttle plasmid, pNEBAd35-2, generating pNEBAd35CMVSEAPBGHpA. The SEAP expression cassette was obtained from a previously constructed shuttle plasmid by *Eco*RI digestion. Following the digestion the desired fragment was gel purified, treated with Klenow to obtain blunt ends and cloned into the *Swa*I site in pNEBAd35-2. The transgene was then recombined into the pAd34ΔE1ΔE4Ad5Orf6, generating pAd34ΔE1SEAPΔE4Ad5Orf6 as described above for the gag transgene.

All pre-Ad plasmids were rescued into virus and expanded to prepare CsCl purified stocks as described above.

#### 10 EXAMPLE 21

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### Construction of pMRKAd34ΔE1ΔE4Ad5Orf6

To construct an Ad34 pre-Ad plasmid that was composed entirely of Ad34 sequences, an Ad34 ITR cassette was generated. The Ad34 ITR cassette was constructed containing sequences from the right (bp 31584 to 31895 and bp 34409 to 34772) and left (bp 4 to 456 and bp 3402 to 3885) end of the Ad34 genome (see Figures 28A-1 to 28A-9) separated by 15 plasmid sequences containing a bacterial origin of replication and an ampicillin resistance gene. These four segments were generated by PCR and cloned sequentially into pNEB193, generating pNEBAd34-4. Next the Ad5 Orf6 open reading frame was generated by PCR and cloned between Ad34 bp 31895 and 34409 generating pNEBAd34-4Ad5Orf6 (the ITR cassette). PNEB193 is a commonly used commercially available cloning plasmid (New England Biolabs 20 cat# N3051S) containing a bacterial origin of replication, ampicillin resistance gene and a multiple cloning site into which the PCR products were introduced. The ITR cassette contains a deletion of E1 sequences from Ad34 bp 457 to 3401 with a unique Swa I restriction site located in the deletion and an E4 deletion from Ad34 bp 31896 to 34408 into which Ad5 Orf6 was introduced in an E4 parallel orientation. In this construct Ad5Orf6 expression is driven by the 25 Ad34 E4 promoter. The Ad34 sequences (bp 31584 to 31895 and bp 3402 to 3885) in the ITR cassette provided regions of homology with the purified Ad34 viral DNA in which bacterial recombination could occur following cotransformation into BJ 5183 bacteria (Figure 27). The ITR cassette was also designed to contain unique restriction enzyme sites (PmeI) located at the end of the viral ITR's so that digestion would release the recombinant Ad34 genome from the 30 plasmid sequences. Potential clones were screened by restriction analysis and one clone was selected as pMRKAd34 $\Delta$ E1 $\Delta$ E4Ad5Orf6.

# EXAMPLE 22 In Vivo Studies

#### A. Immunization

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Cohorts of 3 rhesus macaques were given single intramuscular injections of one of the two vectors: (1) 10^11 vp MRKAd5-SEAP (in MRKAd vector backbone disclosed in PCT/US01/28861, published March 21, 2002); and (2) 10^11 vp Ad34ΔE1SEAPΔE4Ad5Orf6. Rhesus macaques were between 3-10 kg in weight. In all cases, the total dose of each vaccine was suspended in 1 mL of buffer. The macaques were anesthetized (ketamine/xylazine) and the vaccines were delivered i.m. in 0.5-mL aliquots into both deltoid muscles using tuberculin syringes (Becton-Dickinson, Franklin Lakes, NJ). Peripheral blood mononuclear cells (PBMC) were prepared from blood samples collected at several time points during the immunization regimen. All animal care and treatment were in accordance with standards approved by the Institutional Animal Care and Use Committee according to the principles set forth in the Guide for Care and Use of Laboratory Animals, Institute of Laboratory Animal Resources, National Research Council.

#### B. SEAP Assay

Serum samples were analyzed for circulating human secreted alkaline phosphatase (SEAP) levels using TROPIX phospha-light chemiluminescent kit (Applied Biosystems Inc). Duplicate 5 µL aliquots of each serum were mixed with 45 µL of kit-supplied dilution buffer in a 96-well white DYNEX plate. Serially diluted solutions of a human placental alkaline phosphatase (Catalog no. M5905, Sigma, St. Louis, MO) in 10% naïve monkey serum served to provide the standard curve. Endogenous SEAP activity in the samples was inactivated by heating the well for 30 minutes at 65 °C. Enzymatic SEAP activities in the samples were determined following the procedures described in the kit. Chemiluminescence readings (in relative light units) were recorded using DYNEX luminometer. RLU readings were converted to ng/mL SEAP using a log-log regression analyses.

#### C. ELISPOT Assay

The IFN- $\gamma$  ELISPOT assays for rhesus macaques were conducted following a previously described protocol (Allen *et al.*, 2001 *J. Virol.* 75(2):738-749), with some modifications. For antigen-specific stimulation, a peptide pool was prepared from 20-aa peptides that encompass the entire HIV-1 gag sequence with 10-aa overlaps (Synpep Corp., Dublin, CA). To each well, 50  $\mu$ L of 2-4 x 10<sup>5</sup> peripheral blood mononuclear cells (PBMCs) were added; the cells were counted using Beckman Coulter Z2 particle analyzer with a lower

size cut-off set at 80 femtoliters ("fL"). Either 50  $\mu$ L of media or the gag peptide pool at 8  $\mu$ g/mL concentration per peptide was added to the PBMC. The samples were incubated at 37°C, 5% CO<sub>2</sub> for 20-24 hrs. Spots were developed accordingly and the plates were processed using custom-built imager and automatic counting subroutine based on the ImagePro platform (Silver Spring, MD); the counts were normalized to  $10^6$  cell input.

# D. Intracellular Cytokine Staining (ICS)

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To 1 ml of 2 x 106 PBMC/mL in complete RPMI media (in 17x100mm round bottom polypropylene tubes (Sarstedt, Newton, NC)), anti-hCD28 (clone L293, Becton-Dickinson) and anti-hCD49d (clone L25, Becton-Dickinson) monoclonal antibodies were added to a final concentration of 1  $\mu$ g/mL. For gag-specific stimulation, 10  $\mu$ L of the peptide pool (at 0.4 mg/mL per peptide) were added. The tubes were incubated at 37 °C for 1 hr., after which 20 μL of 5 mg/mL of brefeldin A (Sigma) were added. The cells were incubated for 16 hr at 37 °C, 5% CO<sub>2</sub>, 90% humidity. 4 mL cold PBS/2%FBS were added to each tube and the cells were pelleted for 10 min at 1200 rpm. The cells were re-suspended in PBS/2%FBS and stained (30 min, 4 °C) for surface markers using several fluorescent-tagged mAbs: 20 µL per tube antihCD3-APC, clone FN-18 (Biosource); 20 μL anti-hCD8-PerCP, clone SK1 (Becton Dickinson); and 20 µL anti-hCD4-PE, clone SK3 (Becton Dickinson). Sample handling from this stage was conducted in the dark. The cells were washed and incubated in 750 µL 1xFACS Perm buffer (Becton Dickinson) for 10 min at room temperature. The cells were pelleted and re-suspended in PBS/2%FBS and 0.1 μg of FITC-anti-hIFN-γ, clone MD-1 (Biosource) was added. After 30 min incubation, the cells were washed and re-suspended in PBS. Samples were analyzed using all four color channels of the Becton Dickinson FACSCalibur instrument. To analyze the data, the low side- and forward-scatter lymphocyte population was initially gated; a common fluorescence cut-off for cytokine-positive events was used for both CD4<sup>+</sup> and CD8<sup>+</sup> populations, and for both mock and gag-peptide reaction tubes of a sample.

#### E. Results

Expression: Serum samples prior to and after the injection were analyzed for circulating SEAP activities and the results are shown in Figure 29. Results indicate that the peak levels of SEAP protein produced by the alternative adenovirus serotype were lower than but were within 3-fold of that of MRKAd5 at the same high dose level of 10^11 vp (Figure 29). The levels of SEAP in the serum dropped dramatically after day 10 and were close to background as early as day 15. These observations strongly indicate that the Ad34-based vector is efficient in expressing a transgene following intramuscular administration in a primate.

Immunogenicity: Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 30; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells immediately after a single dose of the vector. The responses improved following a second dose given at wk 4. Overall, the responses to the Ad34-based vector were slightly lower than those induced by the same dose of MRKAd5-gag. The results strongly indicate the Ad34-based vector can prime effectively for HIV-specific T cell responses.

IFN-γ ICS analyses of the PBMC from the Ad34-immunized animals revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 31).

#### 15 EXAMPLE 23

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#### Heterologous Immunization

Cohorts of 3 monkeys were immunized (at wks 0, 4) with 10^11 vp Ad34\Delta E1gag\Delta E4Ad5Orf6 followed by a booster at week 24 with 10^10 vp Ad35\Delta E1gag\Delta E4Ad5Orf6. Vaccine-induced T cell responses against HIV-1 gag were quantified using IFN-gamma ELISPOT assay against a pool of 20-aa peptides that encompassed the entire protein sequence. The results are shown in Figure 32; they are expressed as the number of spot-forming cells (SFC) per million peripheral blood mononuclear cells (PBMCs) that responded to the peptide pool or the mock (no peptide) control.

Immunization with gag-expressing Ad34 vector induced detectable levels of circulating gag-specific T cells that decreased to between 94-139 SFC/10^6 PBMC at the time of the boost. Heterologous immunization with an Ad35-based HIV vector resulted in as much as a 3-fold increase in T cell responses.

IFN-γ ICS analyses of the PBMCs from the Ad34 primed/Ad35 boosted animals at week 28 revealed that the vector can induce detectable levels of both CD4<sup>+</sup> and CD8<sup>+</sup> HIV-specific T cells (Figure 33).

#### WHAT IS CLAIMED IS:

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1. A means for propagating replication-defective adenovirus in an adenoviral E1-complementing cell line expressing E1 gene product(s) which are non-native to the adenovirus, which comprises:

- (a) inserting all or a portion of a heterologous adenoviral E4 region comprising nucleic acid sequence encoding open reading frame 6 (ORF6) into a replication-defective adenovirus; wherein the E4 region or portion thereof inserted into the adenovirus is native to a virus of the same adenovirus serotype as the E1 gene product(s) expressed by the complementing cell line;
- (b) introducing the replication-defective adenovirus into the adenoviral E1-complementing cell line;
- (c) allowing the replication-defective adenovirus to propagate in the adenoviral E1-complementing cell line; and
  - (d) rescuing the propagated adenovirus.
- 2. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region.
- 3. A means in accordance with claim 2 wherein the heterologous adenoviral E4 region or portion thereof comprises the complete adenoviral E4-encoding region and native E4 promoter.
- 4. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding open reading frame 6 (ORF6).

5. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is inserted into the replication-defective virus in place of nucleic acid sequence encoding the complete adenoviral E4-encoding region.

- 6. A means in accordance with claim 1 wherein the heterologous adenoviral
   5 E4 region or portion thereof is derived from a subgroup C adenovirus.
  - 7. A means in accordance with claim 1 wherein the subgroup C adenovirus is adenovirus of serotype 5.
  - 8. A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of subgroup B.
  - 9. A means in accordance with claim 7 wherein the replication-defective adenovirus is an adenovirus of serotype 35.

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- 10. A means in accordance with claim 1 wherein the heterologous adenoviral E4 region or portion thereof is operatively linked to a heterologous promoter.
- 11. A means in accordance with claim 1 wherein the adenoviral E1-complementing cell line is a PER.C6® cell line.
  - 12. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6).
  - 13. A replication-defective adenovirus in accordance with claim 12 wherein the adenovirus comprises a heterologous gene of interest.
  - 14. A replication-defective adenovirus in accordance with claim 13 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
  - 15. A replication-defective adenovirus in accordance with claim 14 wherein the HIV-1 antigen is selected from the group consisting of HIV-1 gag, pol, nef and env.

16. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

- 17. A replication-defective adenovirus comprising all or a portion of a

  beterologous E4 region comprising a heterologous adenoviral open reading frame 6 (ORF6) in

  place of a native E4 region or portion thereof comprising ORF6.
  - 18. A replication-defective adenovirus comprising all or a portion of a heterologous E4 region comprising a complete heterologous E4 region in place of a complete native E4 region.
- 19. A replication-defective adenovirus comprising a heterologous E4 region or portion thereof comprising a complete heterologous E4 region including E4 promoter in place of a complete native E4 region.
  - 20. Adenovirus propagated in accordance with the means of claim 1.
  - 21. A means in accordance with claim 1 wherein the replication-defective adenovirus comprises a heterologous gene of interest.

- 22. A means in accordance with claim 21 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.
- 23. A means in accordance with claim 22 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.
- 24. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a heterologous gene of interest.
  - 25. A replication-defective adenovirus in accordance with claim 24 wherein the heterologous gene of interest is a gene encoding an HIV-1 antigen.

26. A replication-defective adenovirus in accordance with claim 25 wherein the HIV-1 antigen is selected from the group consisting of: HIV-1 gag, pol, nef and env.

27. A replication-defective adenovirus of serotype 35 comprising all or a portion of an adenovirus serotype 5 E4 region comprising open reading frame 6 (ORF6) and a gene encoding HIV-1 gag.

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- 28. A recombinant adenoviral vector of serotype 24 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 29. A population of cells comprising the recombinant adenoviral vector of claim 28.
  - 30. A method for producing recombinant, replication-defective adenovirus particles comprising:
  - (a) introducing a recombinant adenoviral vector of claim 28 into a population of cells expressing adenovirus E1; and
    - (b) harvesting the resultant recombinant, replication-defective adenovirus.
  - 31. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 30.
  - 32. A composition comprising purified recombinant adenovirus particles in accordance with claim 31.
  - 33. A composition in accordance with claim 32 which comprises a physiologically acceptable carrier.
  - 34. A recombinant adenoviral vector in accordance with claim 28 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

35. A composition comprising purified recombinant adenoviral particles in accordance with claim 31 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

36. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 35 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 37. A method in accordance with claim 36 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 38. A composition in accordance with claim 35 wherein the heterologous nucleic acid encodes an HTV antigen.
- 39. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 38.
- 40. A composition in accordance with claim 39 wherein the HIV antigen is

  HIV-1 gag or immunologically relevant modification thereof.
  - 41. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
  - 42. A composition in accordance with claim 39 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
  - 43. A recombinant adenoviral vector of serotype 24 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

44. A population of cells comprising the recombinant adenoviral vector of claim 43.

- 45. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 43 into a population of cells expressing adenovirus serotype 5 E1; and

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- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 46. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 45.
- 47. A composition comprising purified recombinant adenovirus particles in accordance with claim 46.
- 48. A composition in accordance with claim 47 which comprises a physiologically acceptable carrier.
- 49. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 48 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
- 50. A method in accordance with claim 49 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 51. A composition in accordance with claim 48 wherein the heterologous nucleic acid encodes an HIV antigen.
- 52. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 51.

53. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 54. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 55. A composition in accordance with claim 51 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

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- A recombinant adenoviral vector of serotype 34 which comprises an E4 gene or a segment of an E4 gene comprising open reading frame 6 ("ORF6") of an alternative serotype.
- 57. A population of cells comprising the recombinant adenoviral vector of claim 56.
- 58. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 56 into a population of cells expressing adenovirus E1; and
  - (b) harvesting the resultant recombinant, replication-defective adenovirus.
  - 59. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 58.
  - 60. A composition comprising purified recombinant adenovirus particles in accordance with claim 59.
    - 61. A composition in accordance with claim 60 which comprises a physiologically acceptable carrier.
    - 62. A recombinant adenoviral vector in accordance with claim 56 which is at least partially deleted in E1 and devoid of E1 activity and comprises a heterologous nucleic acid.

63. A composition comprising purified recombinant adenoviral particles in accordance with claim 59 which are at least partially deleted in E1 and devoid of E1 activity and comprise a heterologous nucleic acid.

64. A method for effecting the delivery and expression of heterologous nucleic acid comprising administering the composition of claim 63 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.

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- 65. A method in accordance with claim 64 wherein the composition is preceded or followed by administration of heterologous nucleic acid with an adenovirus of a different serotype.
- 66. A composition in accordance with claim 63 wherein the heterologous nucleic acid encodes an HIV antigen.
- 67. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 66.
- 68. A composition in accordance with claim 67 wherein the HIV antigen is
  HIV-1 gag or immunologically relevant modification thereof.
  - 69. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
  - 70. A composition in accordance with claim 67 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.
  - 71. A recombinant adenoviral vector of serotype 34 which is at least partially deleted in E1 and devoid of E1 activity; wherein said vector comprises an E4 gene or a segment of an E4 gene from adenovirus serotype 5 comprising open reading frame 6 ("ORF6"), and a heterologous nucleic acid.

72. A population of cells comprising the recombinant adenoviral vector of claim 71.

- 73. A method for producing recombinant, replication-defective adenovirus particles comprising:
- (a) introducing a recombinant adenoviral vector of claim 71 into a population of cells expressing adenovirus serotype 5 E1; and

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- (b) harvesting the resultant recombinant, replication-defective adenovirus.
- 74. Purified recombinant, replication-defective adenovirus particles harvested in accordance with the method of claim 73.
- 75. A composition comprising purified recombinant adenovirus particles in accordance with claim 74.
- 76. A composition in accordance with claim 75 which comprises a physiologically acceptable carrier.
- 77. A method for effecting the delivery and expression of the heterologous nucleic acid comprising administering the composition of claim 76 prior or subsequent to administration of the heterologous nucleic acid with the same or different vector.
  - 78. A method in accordance with claim 77 above wherein the composition is preceded or followed by administration of the heterologous nucleic acid with an adenovirus of a different serotype.
- 79. A composition in accordance with claim 76 wherein the heterologous nucleic acid encodes an HIV antigen.
- 80. A method for generating a cellular-mediated immune response against HIV in an individual comprising administering to the individual a composition of claim 79.

81. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 gag or immunologically relevant modification thereof.

- 82. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 nef or immunologically relevant modification thereof.
- 83. A composition in accordance with claim 79 wherein the HIV antigen is HIV-1 pol or immunologically relevant modification thereof.

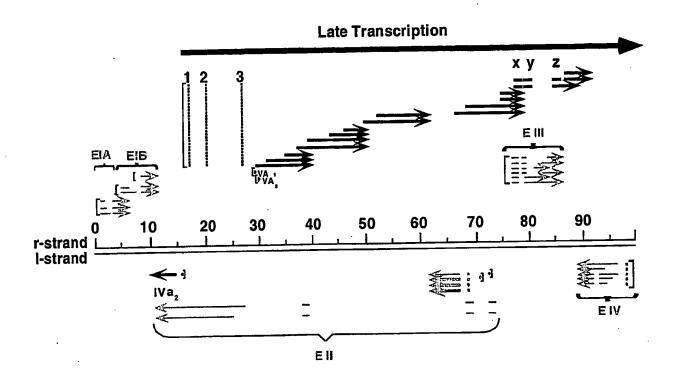


FIG. 1

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42.	andartacca	totcasagto	rectatttt	acotaggigi	cagetyatey	Ctagggtate
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198	1 agaacatgga	aggttcgcaa	gatgaggaca	atettaggtt	actiggicage	gcagcccccg
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210	1 caacacca	acccdadadc	coocctooac	cctccagtgg	aggaggcyya	grayeryace
216	1 tatatacta	actocaacoo	gtgcttactg	gatctacgtc	cactggacgg	gataggggcg
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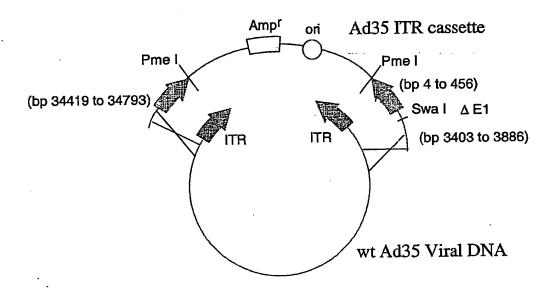


FIG. 3

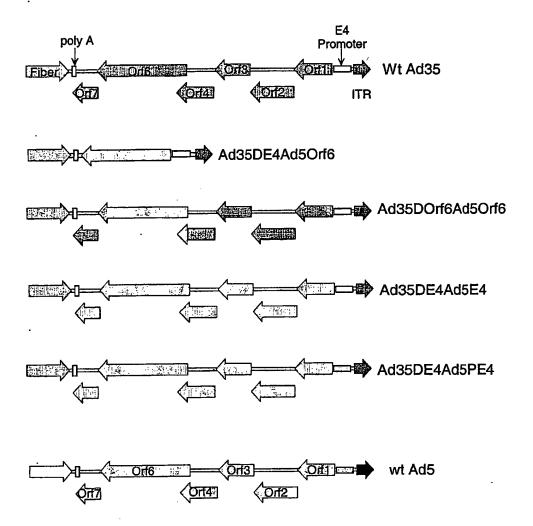


FIG. 4

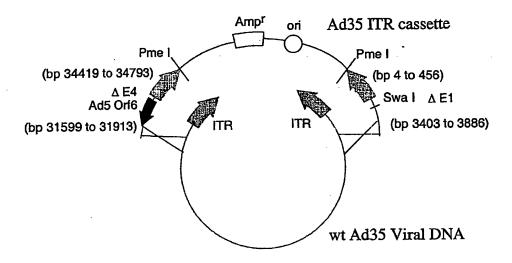


FIG. 5

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1921	CACCATCATG	ATGCAGAGGG	GCAACTTCAG	GAACCAGAGG	AAGACAGTGA	AGTGCTTCAA
1981	CTGTGGCAAG	GTGGGCCACA	TTGCCAAGAA	CTGTAGGGCC	CCCAGGAAGA	AGGGCTGCTG
2041	GAAGTGTGGC	AAGGAGGGCC	ACCAGATGAA	GGACTGCAAT	GAGAGGCAGG	CCAACTTCCT
2101	GGGCAAAATC	TGGCCCTCCC	ACAAGGGCAG	GCCTGGCAAC	TICCICCAGI	CCAGGCCTGA
2161	GCCCACAGCC	CCTCCCGAGG	AGTCCTTCAG	GTTTGGGGAG	GAGAAGACCA	CCCCCAGCCA
2221	GAAGCAGGAG	CCCATTGACA	AGGAGCTGTA	CCCCTGGCC	TCCCTGAGGT	CCCTGTTTGG
2281	CAACGACCCC	TCCTCCCAGT	<i>AA</i> aataaagc	ccgggcagat	ctgatctgct	gtgccttcta
2341	gttgccagcc	atctgttgtt	tgcccctccc	ccgtgccttc	cttgaccctg	gaaggtgcca
2401	ctcccactgt	cctttcctaa	taaaatgagg	aaattgcatc	gcattgtctg	agtaggtgtc
2461	attctattct	ggggggtggg	grggggcagc	acagcaaggg	yyayyattgg	gaagacaata
2521	gcaggcatgc	tggggatgcg	grgggctcta			

SEQ ID NO: 2

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# 16/59

	•					
<u>1</u> _	ccattgcata	cgttgtatcc	atatcataat	atgtacattt	atattggctc	atgtccaaca
61	ttaccgccat	gttgacattg	<u>attattgact</u>	agttattaat	agtaatcaat	tacggggtca
121	ttagttcata	gcccatatat	ggagttccgc	gttacataac	ttacggtaaa	taggeeegeet
181	ggctgaccgc	ccaacgaccc	ccgcccattg	acgtcaataa	tgacgtatgt	teccatagea
241	acgccaatag	ggactttcca	ttgacgtcaa	tgggtggagt	atttacggta	aactgcccac
301	ttggcagtac	atcaagtgta	tcatatgcca	agtacgcccc	ctattgacgt	caatgacggt
361	asataaccca	cctggcatta	tocccaotac	atgaccttat	gggactttcc	tacttggcag
421	tacatctaco	tattagtgat	coctattacc	atogtgatgc	ggttttggca	gtacatcaat
481	gggcgtggat	agcggtttga	ctcacgggga	tttccaagtc	tccaccccat	tgacgtcaat
541	gggagtttgt	tttggcacca	aaatcaacgg	gactttccaa	aatgtcgtaa	caacteegee
601	ccattgacgc	aaataggggg	taggcgtgta	caataaaagg	tctatataag	cagagetegt
661	ttagtgaacc	gtcagatcgc	ctggagacgc	catccacgct	gttttgacct	ccatagaaga
721	ascadadece.	materageet.	ccacaaccaa	gaacggtgca	ttggaacgcg	gatteecege
781	gccaagagtg	<u>agatcgatct</u>	aagtaagctt	CCTGCATGCT	GCTGCTGCTG	CIGCIGCIGG
0/1	CCCTCACCCT	ACACCTCTCC	CTGGGCATCA	TCCCAGTTGA	GGAGGAGAAC	CCGGACTICT
901	GGAACCGCGA	GGCAGCCGAG	GCCCTGGGTG	CCGCCAAGAA	GCTGCAGCC'I'	GCACAGACAG
961	CCGCCAAGAA	CCTCATCATC	TTCCTGGGCG	ATGGGATGGG	GGTGTCTACG	GTGACAGCTG
1021	CCAGGATCCT	AAAAGGGCAG	AAGAAGGACA	AACTGGGGCC	TGAGATACCC	CTGGCCATGG
1081	ACCGCTTCCC	ATATGTGGCT	CTGTCCAAGA	CATACAATGT	AGACAAACAT	GTGCCAGACA
1141	GTGGAGCCAC	AGCCACGGCC	TACCTGTGCG	GGGTCAAGGG	CAACTTCCAG	ACCATIGGCT
1201	TGAGTGCAGC	CGCCCGCTTT	AACCAGTGCA	ACACGACACG	CGGCAACGAG	GTCATCTCCG
1261	TGATGAATCG	GGCCAAGAAA	GCAGGGAAGT	CAGTGGGAGT	GGTAACCACC	ACACGAGTGC
1321	<u>አርር አርርርር ጥር</u>	GCCAGCCGGC	ACCTACGCCC	ACACGGTGAA	CCGCAACTGG	TACTCGGACG
1321	CCCACCTCCC	TGCCTCCGCC	CGCCAGGAGG	GGTGCCAGGA	CATCGCTACG	CAGCTCATCT
1 / / 1	CCAACATGGA	CATTGACGTG	ATCCTAGGTG	GAGGCCGAAA	GTACATGTTT	CGCATGGGAA
1501	CCCCAGACCC	TGAGTACCCA	GATGACTACA	GCCAAGGTGG	GACCAGGCTG	GACGGGAAGA
1561	አጥሮጥርርጥርር ገ	CCAATCCCTC	GCGAAGCGCC	AGGGTGCCCG	GTATGTGTGG	AACCGCACTG
1621	ACCITCATICCA	GGCTTCCCTG	GACCCGTCTG	TGACCCATCT	CATGGGTCTC	TTTGAGCCTG
1691	CACACATCAA	ATACGAGATC	CACCGAGACT	CCACACTGGA	CCCCTCCCTG	ATGGAGATGA
17/1	CACACCCTCC	CCTGCGCCTG	CTGAGCAGGA	ACCCCCGCGG	CTTCTTCCTC	TTCGTGGAGG
1 2 0 1	CTCCTCCCAT	CGACCATGGT	CATCATGAAA	GCAGGGCTTA	CCGGGCACTG	ACTGAGACGA
1051	ጥር አጥርጥጥርር እ	<u> </u>	GAGAGGGCGG	GCCAGCTCAC	CAGCGAGGAG	GACACGCTGA
1921	CCCጥCCጥC \C	TGCCGACCAC	TCCCACGTCT	' TCTCCTTCGG	AGGCTACCCC	CTGCGAGGGA
1 9 9 1	ርርጥ <u>ርር አ</u> ጥርጥጥ	· CGGGCTGGCC	CCTGGCAAGG	CCCGGGACAG	GAAGGCCTAC	ACGGTCCTCC
2041	ጥልጥልሮርርልልል	CGGTCCAGGC	TATGTGCTCA	AGGACGGCGC	CCGGCCGGAT	GTTACCGAGA
2101	CCCACACCCC	GAGCCCCGAG	TATCGGCAGC	: AGTCAGCAGT	GCCCCTGGAC	GAAGAGACCC
2161	» CCC» CCCC»	CCACCTCCCC	GTGTTCGCGC	GCGGCCCGCA	GGCGCACCTG	GTTCACGGCG
2221	TYCENCENCE	GACCTTCATA	GCGCACGTCA	TGGCCTTCGC	CGCCTGCCTG	GAGCCCTACA
2281	CCCCCTGCGA	CCTGGCGCCC	CCCGCCGGCA	CCACCGACGC	CGCGCACCCG	GGTTAAcccg
2341	taatecccaa	attacttcct	ctactaacca	ggacatcagg	tggcccccgc	tgaattggaa
2401	tegateagaa	ttgatctgat	ctgctgtgcc	: ttctagttgc	cagccatctg	ttgtttgccc
2461	ctcccccata	ccttccttga	ccctggaagg	tgccactccc	actgtccttt	cctaataaaa
2521	tgaggaaatt	gcatcgcatt	gtctgagtag	gtgtcattct	attctggggg	gtggggtggg
2581	gcagcacagc	aagggggagg	attgggaaga	caatagcagg	catgctgggg	atgcggtggg
	ctcta					

SEQ ID NO: 3

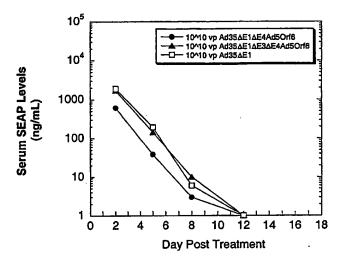
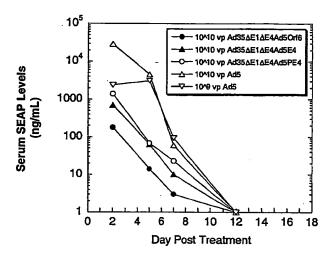


FIG. 8



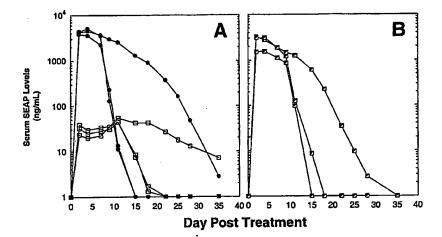


FIG. 10A-B

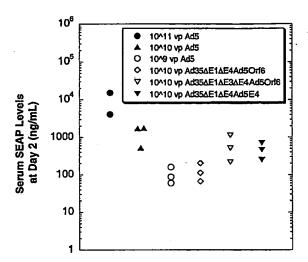


FIG. 11

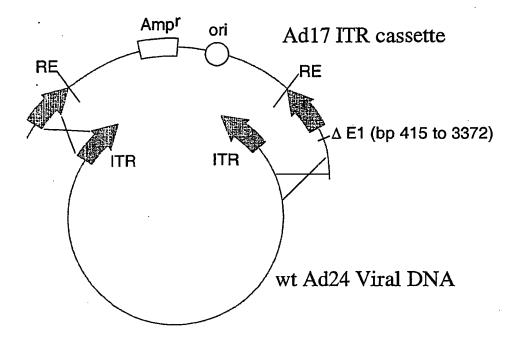
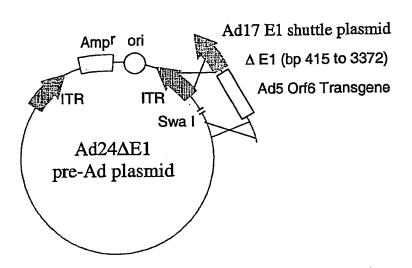


FIG. 12



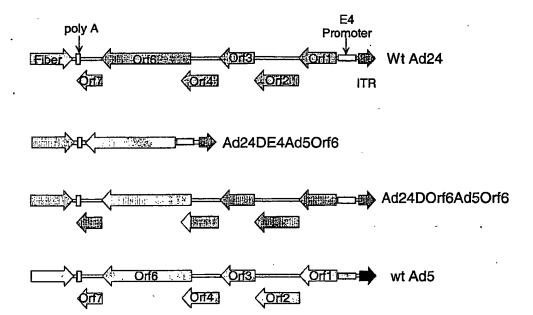


FIG. 14

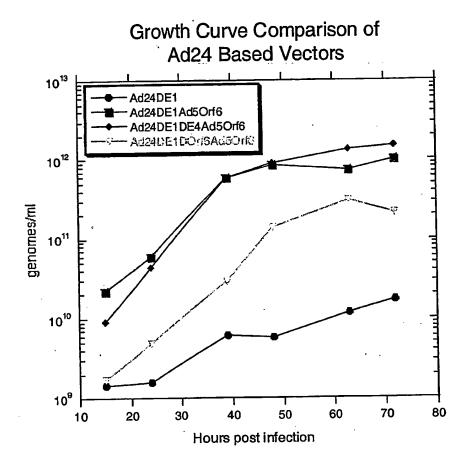


FIG. 15

			agaagtaaa	caaaagttaa	catocaaato	agettttgaa
Τ.	catcatcaat	aatatacccc	acaaaycaaa		tacgcadacg	atcaccacac
9T	tttagggcgg	ggccagcgct	gattygatga	gagaagatga	ageaaatgac	gaactastas
121	acggctaacg	gregeegegg	aggegtggee	tagcccggaa	bbassassass	gggccgacga
181	cgtataaaaa	ageggaettt	agacccggaa	acggccgatt	tteeegegge	cacgcccgga
241	tatgaggtaa	ttctgggcgg	atgcaagtaa	aattaggtca	ttttggegeg	addactydat
301	gaggaagtga	aaagtgaaaa	ataccggtcc	cgcccagggc	ggaatattta	eegagggeeg
361	agagactttg	accgattacg	tgggggtttc	gattgcggtg	ttttttcgcg	aatttccgcg
421	tccgtgtcaa	agtccggtgt	ttatgtcaca	gatcagctga	tccacagggt	atttaaacca
481	gtcgagcccg	tcaagaggcc	actcttgagt	gccagcgagt	agagatttct	ctgagctccg
541	ctcccagagt	ctgagaaaaa	tgagacacct	gcgcctcctt	tcttcaactg	tgcctattga
601	catggccgca	ttattgctgg	aggattatgt	gagtacaata	ttggaggacg	aactgcatcc
661	atctccattt	gagetgggae	ctacacttca	ggacctatat	gatttggagg	tagatgccca
721	tgatgacgac	ccgaacgaag	aggetgtgaa	tttaatattt	ccagaatctc	tgattcttca
781	ggctgacata	gccagcgaag	ctgtacctac	accacttcat	acaccgactc	tgtcacccat
841	acctgaattg	gaagaggagg	acgagctaga	cctccgatgt	tatgaggaag	gttttcctcc
901	cagcgattca	gaggacgaac	agggtgagca	gagcatggct	ctaatctcaa	aatatgcttg
961	tataattata	gaagagcatt	ttgtgttgga	caatcctgag	gtgcccgggc	aaggctgtag
1021	atectoccao	taccaccggg	ataagaccgg	agacacgaac	geeteetgeg	ctctgtgtta
1081	catgaaaaag	aacttcagct	ttatttacag	taagtggagt	gaatgtgaga	gagactgagt
1141	octtaacaca	taactgggta	atocttaaac	agctgtgcta	agtgtggttt	atttttgttt
1201	ctaggtccgg	totcagagga	tgagtcatca	ccctcagaag	aagaccaccc	gtgtccccct
1261	gagetgag	acassacaca	cctgcaagtg	cacagaccca	ccccaqtcaq	acccagtggc
1301	gageegeeag	carctettea	aaaaattgag	gacttgttac	atgacatggg	tggggatgaa
1381	cctttqqacc	tgagettgaa	acccccacc	aactaggctc	agctgtgctt	agtcatgtgt
1//1	anathanatt	ctacaataaa	agtatatoto	acgcatgcaa	agtatagttt	atgactcatg
1601	gaataagtt	actoctatat	asotoocaac	acctgggcac	tagagacacag	accttcaggg
1561	agegeggeee	ageceeatae	actateette	cagactttag	caagacacgc	caacttataa
1601	aguicuigau	agacgcgcgg	tecaaattet	ggagacactg	gtttggaact	cctctatctc
1601	ayyacayccc	cacacttaac	aarrattata	acgaggaatt	tgaaaatctt	tttgctgatt
1001	grerggra	catagetaag	ctasatctcc	gccaccagtc	ccttttccag	gaaagggtac
1001	getetggeet	tentetten	accesace	gcactacagc	coorditact	tttataattt
1001	ttataattaa	caastagaa	cacaacaccc	aactgagcag	ggggtacatt	ctggacttcg
1001	cccggccga	caaatygage	ggaacaccc	ggcagcgggg	acadadaatc	ttgaactact
1001	cagecatyca	cccgcggagg	ccaatcttc	ttcgtctaca	cacacaaaca	tccatgttgg
1301	ggettatata	gccagcagcc	atagacaaa	acccgaggag	caacctaaac	cctccatcaa
2101	aggaagaaat	gaggcaggcc	acgyacgaga	cctgtaccca	gagettagea	gggtgctgac
21.C1	aagaggaget	ggattgaatt	aggiaiccag	gagcgatggg	gageetagea	gggtgctgac
510T	atecatggee	aggggagtga	teasteess	gcgtccagag	cocattacct	ggacgacgac
2221	egagetgacg	gecageerya	cyaattycaa	gatgcaggat	asstatoro	tagaacaaat
2281	acagatggag	tgtagggatg	aggraggere	ttaggaggag	addiategec	aatatoccaa
2341	aaaaacccac	tggttgaacc	cagacyayya	ttgggaggag	accatcaaga	tcacacacac
2401	gatageeetg	cgcccagatt	geaagtacag	ggtgaccaag	acggtgaata	agggaggett
2461	ctgctacatc	teggggaaeg	gggcagaggt	ggtcatcgat	accetygaca	tasttttast
2521	caggtgttgc	atgatgggaa	tgagageegg	agtgatgaat	theateres	agarterest
2581	gaacatgaag	ttcaatggag	agaagtttaa	tggggtgatg	treatgycea	tatagecacae
2641	gaccctgcac	ggctgcagtt	tetteggett	caacaatatg	tgcgcagagg	cetggggege
2701	tgctaagatc	aggggatgta	agttttatgg	ctgctggatg	ggcgtggtcg	yaayacccaa
2761	gagcgagatg	tctgtgaagc	agtgtgtgtt	tgagaaatgc	tacctgggag	tetetaeega
2821	gggcaatgct	agagtgagac	attgctcttc	cctggagacg	ggctgcttct	geetggtgaa
2881	gggcacagcc	tctctgaagc	ataatatggt	gaagggctgc	acggatgagc	gcatgtacaa
2941	catgctgaca	tgcgactcgg	gggtctgcca	tatcctgaag	aacatccatg	tgacctccca
3001	ccccggaag	aagtggccag	tgtttgagaa	taacctactg	atcaagtgcc	acatgcacct
3061	gggcgccaga	aggggcacct	tccagccgta	ccagtgcaac	tttagccaga	ccaagctgct
3121	gctggagaac	gatgccttct	ccagggtgaa	cctgaacggc	atctttgaca	tggatgtctc
3181	ggtgtacaag	atcctgagat	acgatgagac	caagtccagg	gtgcgcgctt	gcgagtgcgg
3241	gggcagacac	accaggatgc	aaccagtggc	cctggatgtg	accgaggagc	tgaggcccga
3301	ccacctggtg	atggcttgta	ccgggaccga	gttcagctcc	agtggggagg	acacagatta
3361	gaggtaggtt	gagtattagt	gggcgtggct	aaggtgacta	taaaggcggg	tgtcttacga
3421	gggtcttttt	gcttttctgc	agacatcatg	aacgggactg	gcggggcctt	cgaagggggg
3481	ctttttagcc	cttatttgac	aacccgcctg	ccgggatggg	ccggagttcg	tcagaatgtg
3541	atgggatcga	cggtggacgg	gcgtccagtg	cttccagcaa	attcctcgac	catgacctac
3601	gcgaccgtgg	ggaactcgtc	gctcgacagc	accgccgcag	ccgcggcagc	cgcagccgcc

FIG. 16A-1

3661	atgacagcga	cgagactggc	ttcgagctac	atgcccagca	gcagcagtag	cccctctgtg
3721	cccagttcca	tcatcgccga	ggagaaactg	ctggccctgc	tggccgagct	ggaagccctg
3781	acccccacc	tggccgccct	gacccagcag	gtgtccgagc	tccgcgaaca	gcagcagcag
3,011	casatasat	gattcaataa	acacagattc	tgattcaaac	agcaaagcat	ctttattatt
2001	tattttta	cgcgcggtag	accetaatee	acctctcccq	atcattgaga	gtgcggtgga
330T	tattttttt	gacccggtag	acctoggett	ggatgttgag	gtacatgggc	atgagcccgt
396T	ttttttccag	gacccggtag	aggraggare	gatgatgag	tagaatcata	ttgtagatga
4021	cccgggggtg	gaggtagcac	cactgcatgg	t-rastrat	atacttasaa	annanactna
4081	tccagtcata	gcaggggcgc	rgggcgrggr	gerggargar	gttcttgagg	aggagaccga
4141	tggccacggg	gagccccttg	grgraggrgr	rggcgaagcg	gitgagetgg	bbeessess
4201	tgcgggggga	gatgatgtgg	agtttggcct	ggatcttgag	grrggcgarg	ttgccaccca
4261	gatcccgcct	ggggttcatg	ttgtgcagga	ccaccagaac	ggtgtagece	gtgcacttgg
4321	ggaacttgtc	atgcaacttg	gaagggaatg	cgtgaaagaa	tttggagacg	cccttgtgcc
4381	cacccaggtt	ttccatgcac	tcatccatga	tgatggcgat	gggcccgtgg	gctgcggctt
4441	toocaaagac	atttctaaaa	tcagagacat	cgtaattatg	ctcctgggtg	agatcatcat
4501	aagacatttt	aatgaatttg	ggggggggg	tgccagattg	ggggacaatg	gttccctcgg
4561	accccaaaac	gaagttcccc	tcacatattt	gcatctccca	ggctttcatc	tcggaggggg
4621	ggatcatotc	cacctgcggg	gcgatgaaaa	aaacggtttc	cggggcgggg	gtgatgagct
4681	осоаооаоао	caggtttctc	aacagctggg	acttgccgca	cccggtcggg	ccgtagatga
1711	ccccatasc	gggttgcagg	tootaottca	aggacatgca	gctgccgtcg	tcccggagga
1801	addadaccsc	ctcgttgagc	atotetetoa	cttggaggtt	ttcccggacg	agctcgccga
4861	ggggggccac	cccgcccagc	gagagcagct	cttqcaqqqa	agcaaagttt	ttcaggggct
4001	taaaccatc	ggccatgggc	atcttggcga	gggtctgcga	gaggagttcg	aggcggtccc
4001	cgagecegee	gacgtgctct	accordatete	gatecageag	acttcctcqt	ttcgggggtt
4901	agageregge	cgactgtagg	acadacaca	ataaacatcc	agcgctgcca	gcgtcatgtc
2041	gggacgactg	ctcagtgtcc	gcacgagacg	actgggcgccc	accotosaco	gatagacccc
5101	cttccagggt	ctcagtgtcc	tagagege	actcatcctc	ctaatactaa	aacaaacaca
5161	gggctgtgcg	cttgcaaggg	tgegettgag	attracecty	aggreege	taaaaacctc
5221	gtcttcgccc	tgcgcgtcgg	cyagatayca	gregaceacg	agecegeage	cadagagaga
5281	ggcggcgtgg	cccttggcgc	ggagettgee	cicggaagag	cycccycagy	cgggacagag
5341	gagggattgc	agggcgtaga	gerraggrae	gagaaagacg	gacttggggg	cgaaagcatc
5401	cgctccgcag	tgggcgcaga	cggtctcgca	ctcgaccagc	caggtgaget	egggetgete.
5461	ggggtcaaaa	accagttttc	ccccgttctt	tttgatgcgc	ttettacete	gegteteeat
5521	gagtctgtgt	ccgcgctcgg	tgacaaacag	gctgtctgtg	teceegtaga	cggacttgat
5581	agacctatcc	tgcagggggg	tecegeggte	ctcctcgtag	agaaactcgg	accactctga
5641	gacgaaggcg	cacatecaca	ccaagacaaa	ggaggccacg	tgcgaggggt	agcggtcgtt
5701	atccaccaga	gggtccacct	tttccacggt	atgcagacac	atgtccccct	cctccgcatc
5761	caagaaggtg	attaacttat	aggtgtaggc	cacgtgaccc	ggggtccccg	acgggggggc
5821	ataaaagggg	acaaatetat	actcatcctc	actctcttcc	gcgtcgctgt	ccacgagcgc
5881	cagetattag	ggtaggtatt	ccctttcgag	agcgggcatg	acctcggcac	tcaggttgtc
5941	agtttctaga	aacgaggagg	atttgatgtt	ggcttgccct	gccgcaatgc	tttttaggag
6001	actttcatco	atctggtcag	aaaagactat	ttttttattg	tcaagcttgg	tggcgaagga
6061	accatagaga	gcgttggaga	gaagettgge	gatggatctc	atggtctgat	ttttgtcacg
6121	gecaeagagg	tccttggccg	cgatgttgag	ctggacatac	tegegegega	cgcacttcca
6101	ttcaaaaaa	acggtggtgc	acteateaga	cacgatecto	acgcgccagc	cgcggttatg
0101	coccegggaag	agatccacgc	taataaccac	ct.caccacac	aggggctcgt	togtccagca
0241	eagggreace	ceettgegeg	accadaacdd	addcadcaca	tcaagcagat	actcatcaga
630T	gaggegreeg	tcgatggtga	ageagaaegg	acadagttco	ttotcaaaat	aatcgatttt
030T	ggggteegea	tcatccaagg	agatgtccgg	ctcacaaaca	accadedete	actegtaggg
6421	tgaggatgca	ggacccaagg	ccatccgcca	catcagaggag	, geodgegeee	taccacagat
6481	gttgagggg	ggaccccagg	gcacgggacg	cgccayggcg	guggegtaea cataaaataac	accaccaca
6541	gtcgtagaca	tagatgggct	ccyayayyat	gccgacgcag	, accaccasaca	agogoooco
6601	gcggatgctg	gegegeaegt	agtcatacaa	cccgcgcgag	ggggccaaga	taggegggee
6661	gagattggtg	cgctggggct	gcccggcgcg	gaagacgacc	: tygtyaaaya	. tggcatgcga
6721	.gttggaggag	atggtgggcc	gttggaagat	gttaaagtgg	, gcatgaggca	. yacyaaccya
6781	gtcgcggatg	aagtgcgcgt	aggagtcttg	cagettggeg	, acgagetegg	cggrgacgag
6841	gacgtccatg	gcgcagtagt	ccagcgtttc	gcggatgatg	rcataacccg	COLOROCTE
6901	cttctcccat	: agctcgcggt	tgagggcgta	ctcctcgtca	tccttccagt	actcccggag
6961	coggaatect	: cgatcgtccg	cacggtaaga	, gcccagcato	, tagaaatggt	tcacggcctt
7021	gtagggacag	r cagcccttct	ccacggggag	, ggcgtaagct	tgagcggcct	: tgcggagcga
7081	aatatacata	agggcgaagg	tatccctgac	: catgactttc	: aagaactggt	. acttgaaatc
7141	caaatcatca	r cageegeegt	gctcccagag	, ctcgaaatc	g gtgcgcttct	: tcgagagggg
7201	gttaggcaga	a gcgaaagtga	cqtcattgaa	a gagaatcttg	g cctgcccgcg	f gcatgaaatt
7261	acagataata	g cggaaaggg	ccggaacaaa	ggctcggtt	ttgatgacct	gggcggcgag
, 202				· •	_	

FIG. 16A-2

2201		tcgaagccgt	tastattata	cccaacaata	tagagttcca	tgaatcgcgg
7321	gacgateteg	tegaageegt	cyatyttyty	stastastas	atasaataat	caaaacaata
7381	geggeettta	atgtgcggca	gctttttgag	Ciccicgiag	gtgaggtttt	tesstesses
7441	cagtccgtgc	tgctcgagcg	cccactcctg	gagatgtggg	ttggcttgca	cgaacyaayc
7501	ccadadctcd	coooccataa	gggtctggag	ctcgtcgcga	aagaggcgga	actgctggcc
7561	cacqqccatc	ttttctagag	tgacgcagta	gaaagtaagg	gggtcccgct	cccagcgatc
7621	ccaccataaa	cgcacggcta	gatcgcgagc	gagggggacc	agctctgggt	ccccgagaa
7021	thackage	agcataaagg	adecdedeta	cttgccgaag	gaccccatcc	aggtgtaggt
7681	tttcataacc	agcalaaayy	ggacgagetg	cotgoogaag	tasasaccas	ttaaaaaaaa
7741	ttctacatcg	taggtgacaa	agageegete	cytycyayya	testesses	2022540000
7801	ctggatttcc	tgccaccagt	tggacgagtg	gctgttgatg	tgatgaaagt	ayaaatcccg
7861	ccggcgaacc	gagcactcgt	gctgatgctt	gtaaaagcgt	ccgcagtact	cgcagcgctg
7921	cacagactat	acctcatcca	caagatacac	agcgcgtccc.	ttgaggagga	acttcaggag
7981	tagagagat	aactaataat	tttcatgttc	gcctgcgtgg	gactcaccct	ggggctcctc
0041	asacsacaaa	aggctgacga	acceacacaa	gagccaggtc	cagatetegg	cgcggcgggg
0141	yayyacyyay	aagacgaggg	cacacaatta	ggaggtgtcc	atggtgtcgc	ggagatccag
STOT	geggagageg	aayacyayyy	cycycagecy	atagaagaaa	atasaaacat	acttaggata
8161	gtccgggggc	agggttctga	ggttgacett	grayaggray	grgagggcgt	accedagees
8221	cagatggtac	ttgatctcca	cgggtgagtt	ggrggcrgrg	Lecaegeatt	gcatgagccc
8281	gtagetgege	agaccacga	ccgtgccgcg	gtgcgctttt	agaagcggtg	regeggaege
8341	acticcaaca	acaacaacaa	ttccggcccc	gcgggcaggg	gcggcagagg	cacgtcggcg
8401	tagegetegg	gcaggtcccg	atactacacc	ctgagagcgc	tggcgtgcgc	gacgacgcgg
9/61	caattaacat	cctggatctg	ceacetetae	gtgaagacca	ccggccccgt	gactttgaac
0501	characas	gttcaacaga	atcaatctcg	gcgtcattga	caacaaccta	acgcaggatc
8521	ctgaaayaca	cgcccgagtt	atcaucctog	gogotatoga	acatgaactg	ctccatctcc
8581	tcttgcacgt	cgcccgagtt	gteetggtag	gegatetegg	acacgaactg	attggaggtg
8641	tcctcctgga	gatcgccgcg	gcccgcgcgc	tccacggtgg	eggegaggee	actygagaty
8701	cgacccatga	gctgcgagaa	ggcgcccagg	ccgctctcat	tccagacgcg	getgtagace
8761	acatececat	caacatcaca	cacacacata	accacctgcg	cgaggttgag	ctccacgtgc
8821	cacataaaaa	caacataatt	gcgcaggcgc	tggaagaggt	agtttagggt	ggtggcgatg
8881	tactcaataa	cgaagaagta	catgatccag	cggcgcaggg	gcatctcgct	gatgtcgccg
0001	staggega	gcctttccat	aacctcataa	aaatccacag	cgaagttgaa	aaactgggcg
0741	atgyceteca	agaccgtgag	ggootogtag	annancetna	tgagttcggc	gatggtggcg
900I	rrgegggeeg	agaccgrgag	cccgccccc	tostactatt	cctcttcttc	catgacgacc
9061	cgcacctcgc	gctcgaaatc	cccgggggcc			24434444
9121	tcttcttcta	tttcttcctc	tgggggcggt	garaaraaca	gggcccgacg	acyacygcya
9181	cacaccaaaa	gacggtcgac	gaagegeteg	atcatctccc	cgcggcggcg	acgcatggtt
9241	traataacaa	cacaacccca	ttcqcqagga	cgcagcgtga	agacgccgcc	ggtcatctcc
9301	contaatooo	acaaatcccc	attaggcagc	gagagggcgc	tgacgatgca	tcttatcaat
0261	tacaatataa	gggacgtgag	cacatcaaaa	tcgaccggat	cggagaatct	ttcgaggaaa
0421	rectetare	aatcgcagtc	acaaaataaa	ctcaaacacg	tagcagccct	gtggacgctg
9421	gegeetagee	ggttgctgat	geauggeaug	aadtaddcat	ttttaaggcg	acagatagta
9481	ttagaattge	ggttgctgat	yatytaatty	Lastasstas	gaaggag	gaggatggag
9541	gcgaggagga	ccaggtcctt	gggteeeget	tgetggatge	gaageegeee	agtetestes
9601	caggcctggc	cctgacaccg	gctcaggttc	ttgtagtagt	catgcatgag	ceteteaatg
9661	tcatcactgg	cggaggcgga	gtcttccatg	cgggtgaccc	cgacgcccct	gageggetge
9721	acqaqcqcca	agtcagcaac	gacgcgctcg	gcgaggatgg	cctgttgcac	gcgggtgagg
9781	gtgtcctgga	agtcgtccat	gtcgacgaag	cggtggtagg	ccccggtgtt	gatggtgtag
00/1	ataceattaa	ccatgagcga	ccagttgacg	gtctgcaggc	cgggttgcac	gacctctgag
0001	tocatacas	gcgagaaggc	acacaaatca	aagacatagt	cottocaggt	gcgcacgagg
3301	taccigagec	caactaggaa	atacaacaac	aactaacaat	adadcddcca	acactagata
9961	tactggtate	caactaggaa	gtgtggtggt	approgramme at the second	agagoggota	dadatadcad
10021	gccggcgcgc	ccggggccag	greeregage	acyaggegge	ggcagccgca	gaggaagtga
10081	gacatccagg	tgatgccggc	ggcggtggtg	gaggegegeg	ggaactegeg	gacgeggeee
10141	cagatgttgc	gcagcggcag	gaaatagtcc	atggtcggca	eggtetggee	ggtgagacgc
10201	gegeagteat	tgacgctcta	gaggcaaaaa	cgaaagcggt	tgagcgggct	cttcctccgt
10261	agectggegg	aacocaaaco	ggttaggccg	cgtgtgtacc	ccggttcgag	tecectegaa
10201	teagactage	gccgcgacta	acgtggtatt	ggcactcccg	tctcgacccg	agcccgatag
10321	ccaggergga	acggcggaga	accetttta	ccdaccdadd	ggagtcgcta	gacttgaaag
10381	ccgccayyat	acggcggaga	tostsss	ccgaccgagg	ctagaaaaac	tttaccaaaa
10441	cggccgaaaa	ccccgccggg	tagtggeteg	cyceegtage	ccggagaage	actoocaggg
10501	ttgagtcgcg	gcagaacccg	gttcgcggac	ggccgcggcg	aycygyactt	ggccaccccg
10561	ccgatttaaa	gacccacage	cagccgactt	ctccagttac	gggagcgagc	CCCCCCCCCC
10621	ctttttgcca	gatgcatccc	gteetgegee	aaatgcgtcc	caccccccct	ccggcgacca
10681	ccacaaccac	gaccataaca	agcaccaaca	ctgtagcccc	gccacagcag	acagagatgg
10741	acttonaana	gggcgaaggg	ctggcgagac	tgggggcacc	gtccccggag	cgacaccccc
10001	~~~	gcagaaggac	atacacccaa	catacataca	tacacagaac	ctattcaaaa
TOROT	gegegeaget	gcayaayyac	gogogoogg	acaactacaa	ttttcaaaca	addagagaaa
10861	accgcagcgg	ggaggagccc	gaggagatge	######################################	ccccgggcg	220232322
10921	tgcgcgaggg	cctggaccgc	cagegegege	Lycycyacya	gyacttegay	ccyaacyayt

FIG. 16A-3

10981	agacggggat	cagccccgcg	cgcgcgcacg	tggcggcggc	caacctggtg	acggcctacg
110/1	accacaccoot	gaaggaggag -	cacaacttcc	aaaagagttt	Caacaaccat	gracacac
11101	taatcococo	caaaaaaata -	accctaaact	tgatgcacct	grgggaccrg	gcggaggcca
11161	taatagagag	cccggacagc	aagcctctga	caacacaact	gttcctggtg	gtgcagcaca
11221	ccgcgcagaa	cgaggcgttc	addaaddcac	toctaaacat	cgccgagccc	gagggccgct
11221	geagggaeaa	gctgatcaac	atettecaga	gcatcgtagt	gcaggagcgc	agcctgagcc
11781	ggetgetgga	ggtggcggct	atcocycaga	contactasa	cctgggcaag	ttttacgcgc
11341	tggccgagaa	ggtggegget	taggtgggg	taracaarra	ggtgaagata	gacagetttt
11401	gcaagattta	caagacgccg	caegugeeea	tagacaagga	cctaaacata	taccgcaacg
11461	acatgcgcat	ggcgctcaag	gtgetgaege	tgagegaega	cctgggcgcg	cacagoatas
11521	accgcatcca	caaggccgtg	agegegagee	ggeggegega	gergagegae	toctacttco
11581	tgctgagtct	gcgccgggcg	ctggtagggg	gegeegeegg	cggrgaggag	gootaccuts
11641	acatgggggc	ggacctgcat	tggcagccga	gccggcgcgc	ettggaggee	gcccacggcc
11701	cagaggactt	ggatgaggat	gaggaagagg	aggaggatgc	accegetgeg	gggtactgac
11761	acctccataa	tototttta	gatgcagcaa	gccccggacc	ccgccataay	ggcggcgccg
11921	casagccagc	catecaatet	agcatcggac	qactgggagg	ccgcgatgca	acgcarcary
11221	accetaacaa	cccccaaccc	cgagtccttt	agacaacagc	cgcaggccaa	cagacteteg
119/1	accattetaa	aggcggtggt	cccctctcgg	accaacccca	cgcacgagaa	ggrgerggeg
12001	atcotoaaco	cactaacaa	gaacaaggcc	atccgtcccg	acgaggccgg	gergergrae
12061	aacgccctgc	tagagggggt	agaccactac	aacagcacaa	acgtgcagtc	caacciggac
12121	caactaataa	caracataca	caaaaccata	acacaacaca	agcggttcaa	gaacgagggc
12181	ctaggetegt.	taataacact	gaacgccttc	ctggcgacgc	agccggcgaa	egrgeegege
122/1	acacacaca	attacaccaa	ctttatcagc	acactacaac	tgatggtgac	cgaggtgccc
10201	gggcaggacg	tgtaccagtc	gggcccagac	tactttttcc	agacgagccg	gcagggcttg
12261	cagagegagg	acctaagcca	ggctttcaag	aatctgcgcg	gactatagag	cgtgcaggcg
T730T	cagacggrga	accggtcgac	ggtcatcag	ttgctaacgc	ccaactcgcg	gctgctgctg
12421	ceegragaea	cgcccttcac	cuacaucuuc	agcgtgaacc	gcaactcgta	cctgggccac
T248T	etgetgateg	tttaccgcga	aaccataaac	cadacacada	togacgagca	gacettecag
17241	cigetyacyc	gcgtgagccg	cacactagge	cagaacgaca	ccgacagtct	gagagccacc
12601	gagateaeta	tgctgacaaa	taracarcar	agaattccoo	cacaatacac	actateaace
12661	ctgaacttct	gcatcctgag	Lagacageag	cagaccctag	ggetttteet	gatgcaggag
12721	gaggaggagc	geaceetyay	acacycycay	agagegeag	acatogaacc	taggatgtag
12781	ggggccaccc	ccagcgccgc	gerggacarg	accycycyca	tacaccacac	aactaccata
12841	gccgccaacc	ggccgttcat	caataagety	acgyactact	ageteceaee	accagaatte
12901	aactcggact	actttactaa	tgetataeta	aaccegcact	tactatagae	cascataasc
12961	tacacgggcg	agtacgacat	gcccgacccc	aacgacgggc	caateaace	accacasac
13021	agcgcggtgt	tctccccgac	cttgcaaaag	egecaggagg	cygtacycac	attaccaaac
13081	gagggcgcgg	tgggtcggag	CCCCTTCCT	agettaggga	guilgualag	atacatasac
13141	tcggtgaaca	gcggcagggt	gagccggccg	egettgetgg	gcgaygacga	gracergaac
13201	gactcgctgc	tgcagccgcc	gegggtcaag	aacgccatgy	ccaataacyy	tacagagaga
13261	ctggtggaca	aactgaaccg	ctggaagacc	tacgctcagg	accataggga	racateges
13321	ccgcggcgac	agcgccacga	ccggcagcgg	ggcctggtgt	gggacgacga	ggaeteggee
12281	racratarca	gcatattaga	cttaaacaaa	aqcggtgggg	ccaacccgtt	egegeatety
13///1	carcccarac	t.ggggcgacg	gatgttttga	atgaaataaa	. actcaccaag	gccatagege
13501	acattetett	ccttattaga	gatgagggg	geggtggtgt	CETCCECCC	tectecticg
13561	taccacacco	taataacaca	agcaaccctq	gaggttccgt	ttgtgeetee	geggeacacy
13621	actectacac	agggcagaaa	cagcattcgt	tactcggaac	: tggctccgca	gracyacacc
12601	actededtat	· acttootoga	caacaagtcg	gcggacatcg	cttccctgaa	Claccadae
127/11	- marcarance	. acttcctgae	cacaataata	cagaacaacg	atticacec	cyccyayycc
12001	2002000202	, cσataaattt	taacaaacaa	tcacaatggg	geggtgatti	. gaayaccarr
12061	ctocacacca	acatocccaa	totoaacoao	tacatgttca	ı ccagcaagıı	. taaggegegg
13021	atastaataa	r ctaggaaggt	ggtagatcac	r aatqatagga	ı gcaaggatga	gttaaaatat
12001	gagtggttt	r agtittaccct	acccaaaaa	aacttttccg	, agaccalgac	catagaccig
1/10/11	atmaacaaco	, ccatettqqa	aaactactto	r caaqtgggg	: ggcaaaatgg	, cgrgcrggay
1 / 1 / 1 / 1	accostated	, gagtcaagtt	tgacagcago	, aatttcaago	: tgggctggga	cccggtaacc
1/161	l aagctggtg	tacctagaat	. ctacacctac	gaggccttcc	: acccggacgt	tgrgcrgcrg
1/221	ceanactaca	, gagtagaett	caccgagagg	cacctgage	a acctectggg	, cattegeaay
1/201	l aagraaccti	- tccaagaggg	cttcaggato	: atotatgagg	g atctcgaggg	, tggtaacatc
1/2/1	l cocaccate	- togatotcaa	gcaatattt	r gatagtaaaa	a agaagcttga	a ggaggcaaca
1 4 4 0 1	l cacastoca:	- ccagggctgc	· togagatato	agaggagaca	a gtcatattco	: aagagctgig
1440.	L cayaatyca	g ctgaaaagga	tetaatesti	gtaccagtag	a cacaagato	a aagtaagaga
1440.	. gaacaagegg	g tcatagatgg	. cacccatca	accetetace	gaagttggta	a cctgtcctat
1452.	agctataat	g acceegagaa	, cacciacya	tcatagaca	toctcacca	cccggacgtc
1458	acctacggg	, accordagad	. aaaaaacacat	,		

FIG. 16A-4

		_				
14641	acctgcggcg	cggagcaagt	ctactggtcg	ctgccggacc	tcatgcaaga	ccccgtcacc
14701	ttcccctcta	cccagcaagt	cagcaactac	cccataatta	gcgccgagct	catgcccttc
44701	cccgccca			atataataa		cacctacacc
14/61	cgcgccaaga	gcttttacaa	egacetegee	gictactece	agctcatccg	cayclacacc
14821	tccctcaccc	acgtcttcaa	ccgcttcccc	gacaaccaga	tcctctgccg	tccgcccgcg
1/1001	aaaaaataa	ccacaatcaa	trassacrito	cctactctca	cagatcacgg	gacgctaccg
TAGOT	cccaccacca	ccacggccag				taggggggg
14941	ctgcgcagca	gtatccgcgg	agtccagcga	gtgaccgtca	ctgacgcccg	Legeegeace
15001	totccctaco	tctacaaggc	cctgggcata	gtcgcgccgc	gcgtgctttc	cagtcgcacc
15061	ttataaaaa	tatatattat	catctcccc	agcaataaca	ccggctgggg	tettactagg
12001	ttttaaaaaa	Lyccatte	caccacgeoc			acceptage
15121	cccagcacca	tgtacggagg	agccaagaag	egeteeeage	agcaccccgt	eegeg ceege
15181	ggccacttcc	gegeteeetg	gggcgcttac	aagcgcgggc	ggacttctac	cgccgccgtg
152/1	cacaccacca	tenachaent	categacteg	ataatcacca	acgcgcgcaa	ctataccccc
15241	cycaccaccy	b		a wastastas	000000000	casctataca
15301	gccccctcca	ccgtggacgc	ggtcatcgac	agegraggragg	ccgacgcgcg	cyactatyce
15361	agacgcaaga	gccggcggcg	acggatcgcc	aggcgccacc	ggagtacgcc	cgccatgcgc
15421	accaccaaa	ctctactaca	ccacaccaaa	cacacaaacc	gccgggccat	gatgcgagcc
15401	9009000999	aaaaaaataa	2000000000	adcadastc	gcagacgagc	aaccaccacc
15481	gegegeegeg	cogcoactgc	acceccegea	ggcaggaccc	gcagacgagc	2500500500
15541	gctgccgcgg	ccatttctag	catgaccaga	cccaggcgcg	gaaacgtgta	ergagracae
15601	gactccgtca	caaacataca	catacccata	cgcacccgtc	ctcctcgtcc	ctgatctaat
15661	anttatata	teccecae	acascastat	caaagcgcaa	aatcaaggag	gagatgetee
12001	gerrara	cccccgcaa	gegaegaege		22022222	gagarates
15721	aggtcgtcgc	cccggagatt	tacggaccac	cccaggcgga	ccagaaaccc	Cycaaaacca
15781	agcgggttaa	aaaaaaggat	gaggtggacg	agggggcagt	agagtttgtg	cgcgagttcg
15841	ctccacaaca	acacataaat	tagaaggggc	gcagggtgca	gcgcgtgttg	cggccggca
12041	ccccgcggcg	gegegeaaae		actoratora	gagagagaat	aggtatgecg
12201	cggcggrggr	gtttacgccc	ggcgagcggt	ceteggteag	gagcaagcgt	agctatgacg
15961	aggtgtacgg	cgacgacgac	atcctggacc	aggcggcgga	gcgggcgggc	gagttcgcct
16021	acqqqaaqcq	gtcgcgcgaa	gaggagetga	teteattace	gctggacgag	agcaacccca
16001	acceptace	gaagagaata	accetacaac	anatactacc	ccaagcagtg	ctactaccaa
10091	egeetageet	gaageeegeg	accertigeage	aggreecec	t	-tt
16141	gccgcggggt	caagcgcgag	ggcgagaata	tgtacccgac	catgcagatc	atggtgeeea
16201	agcgccggcg	cqtqqaaqaa	gtgctggaca	ccgtgaaaat	ggatgtggag	cccgaggtca
16261	agatacacca	catcaaccac	ataacaccaa	acctagacat	gcagaccgtg	gacattcaga
10201	aggigegeee	,caccaagcag	gragegeegg	9000999090	godgaoogtg	900000000
16321	tccccaccga	catggatgtt	gacaaaaaac	cetegaceag	catcgaggtg	cagaccgacc
16381	cctggctccc	agcetecace	gctgccgtct	ccacttctac	cgccgccacg	gctaccgagc
16441	ctccagaag	gcgaagatgg	ggccctgcca	accooctoat	gcccaactac	gtattgcatc
10241	ccccagaag	9094494099	acatataca	acadocata	ctacgccagc	cacaaacacc
TOOUT	cttccattat	eeegaegeeg	ggctategeg	gcacceggca	Ctacgccagc	cgcaggcgcc
16561	cagccagcaa	acgccgccgc	cgcaccgcca	cccgccgccg	tctggccccc	gcccgcgtgc
16621	accacataac	cacqcqccqq	ggccgctcgc	tegttetgee	caccgtgcgc	taccacccca
16691	gratactta	atcontatac	totoatacto	ttgcagagag	atggctctca	cttaccacct
10001	gcattetta	accogcgcgc			223322222	taageagaaa
16/41	gcgcatcccc	gtcccgaatt	accgaggaag	acceegeege	aggagaggca	cygcaggcag
16801	cggcctcaac	cgccgccggc	ggcgggccat	gcgcaggcgc	ctgagtggcg	gctttctgcc
16861	cacactcata	cccataatco	cooccoccat	cggcacgatc	ccgggcatag	cttccgttgc
16001		tarangara	attastatas	restasance	tctttagact	ctmacacacc
10921	getgeaggeg	cegeagegee	gitgatgige	gaataaagcc	-tt	
16981	tggtcctgta	tatttttaga	atggaagaca	tcaattttgc	gtccctggct	ccgcggcacg
17041	gcacgcggcc	gttcatgggc	acctggaacg	agatcggcac	cagccagctg	aacgggggcg
17101	cetteaatte	gaggagtgtc	tagagggggg	ttaaaaattt	cggctcgacg	ctccggacct
17101	ccccaacty	gugcugcgcc			22222222	atananana
1,191	atgggaacaa	ggcctggaat	agcagcacgg	ggcagttgtt	aagggaaaag	Cicaaayacc
17221	agaacttcca	gcagaaggtg	gtggacggcc	tagcctcggg	cattaacggg	gtggtggaca
17281	tagcaaacca	aaccatacaa	cocoaoataa	acagccgcct	ggacccgcgg	ccaccacag
17241	tagoaaaooa	22002020	actostoss	cacacasaaa	cgagaagcgg	ccacaaccca
1/341	tggtggayat	ggaagatgta	accectege	cycccaaggg	-bases	cegeggeeeg
17401	acgcggagga	gacgatcctg	caggtggacg	agccgccctc	gtacgaggag	gccgtcaagg
17461	ccaacatacc	caccacgcgt	atcatcgcgc	cactggccac	tggtgtaatg	aaacccgcca
17521	coattasact	acctecacea	cccacaccca	ctccaccaa	ggcagctccg	attatacaac
1/321	ccccigaccc	geeteegeea			554454445	5005050050
17581	cccctcctgt	ggcgaccgcc	gracaceaca	receegeecy	ccgccaggcc	cagaactggc
17641	agagcacgct	gcacagtatc	gtgggcctgg	gagtgaaaag	tctgaagcgc	cgccgatgct
17701	attranarar	aggaaagagg	acactaaagg	gagagettaa	cttgtatgtg	ccttaccccc
17761		-222-25	taccccctcc	atratrongo	agtgggcgta	catocacato
T//0T	agagaacgcg	cyaayacyyc	Lacticities	acyacyccyc	-303330304	
17821	gccgggcagg	acgcctcgga	gtacctgagc	ccgggtctgg	tgcagtttgc	ccgcgccacc
17881	gacacgtact	tcagcctggg	caacaagttt	aggaacccca	cggtggctcc	cacccacgat
17041	atanconce	accuat ccca	acatatasaa	ctacacttta	tgcccgtgga	teacasaase
10000	gryactacyg	interpretation			22-22-22	
18001	accacgtact	cgtacaaggc	yegetteact	crggccgrgg	gcgacaaccg	ygtgctagac
18061	atggccagca	cttactttga	catecgegge	gtcctggacc	gcggtcccag	cttcaaaccc
18121	tactcoooce	coacttacaa	cagectages	сссаваеес	ccccaactc	tagtcagtgg
10121				22222222	acacatttco	5
TRIRT	gaacaagcta	aagctaccaa	Lyccyyttaa	aayyaaactC	acacatttgg	agrageoget
18241	atgggcggag	aagacattac	agtgaaaggt	cttcaaattg	gaactgatga	aactaaggaa

FIG. 16A-5

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						artagaaaa
18301	gatggagagg	atgaaatttt	tgcagatcaa	acattccagc	cagaacctca	ayuyyayaa
18361	cagaactggc	aagaaacgtt	tgttttctat	ggaggcagag	ctcttaagaa	agaaaccaaa
18/21	atgaagccat	ottatooctc	ttatocoaca	cccacaaatq	aaaagggagg	acaggctaaa
10421	tttacacttg	2442233444	tcacccaacc	aaaattccto	atattacaat	ggatttcttt
10401	gatagtccac	atyaaaaayy	-tagectate	addattoctg	gagatattgt	catotatoca
18541	gatagtccac	aagatgatac	accaggigia	actaataayt	cagacaccgc	caegeatgea
18601	gaaaatgtaa	atttagaagc	tcctgacaca	catgtagttt	acaaaccayy	caaayacyac
18661	tctagttctt	ccgctaacct	cacacaacag	gccatgccta	acagaccgaa	ctacateggg
18721	ttcagagaca	actttgtggg	tcttatgtac	tacaatagta	ctggcaacat	gggtgtgctg
18781	gctggtcagg	cctctcagtt	gaatgctgtg	gtcgacttgc	aagacagaaa	caccgagctg
199/1	tcttaccagc	tattoctaga	ttctctagat	gacagaacca	gatactttag	catgtggaat
10041	tctgcagtgg	agaggtatga	ccccatate	aggatcattg	agaatcacgg	tatagaagat
10301	gaacttccaa	acagetacga	accognegat	aggattaatt	ctaacaccac	atacaaaggt
T830T	gaactteeaa	actatigett	Cocactyaat	ggcagtggcc	etattageac	acasastcac
19021	gttaaagctg	gaactggaaa	caattgggat	gacyargaaa	atyctycaay	acaaaaccag
19081	attggcactg	gcaacctgtt	cgccatggag	atcaacctcc	aggecaacet	acygaayayc
19141	tttctgtact	cgaacgtggc	cctgtacctg	cccgactcct	acaagtacac	gccggccaac
19201	atcacactac	ccaccaacac	caacacctac	gactacatga	acggccgcgt	ggtagccccc
19261	teactaataa	acccctacat	caacattggc	accegetagt	cgctggaccc	catggacaat
10371	gtcaatccct	tcaaccacca	ccgcaacgcg	aacctacact	accgctccat	gctcctgggc
10301	aacggccgct	acatacactt	ccacatccaa	gtgcccaaa	agttctttgc	catcaagaac
12301	ctgcttctgc	teeegette	ctacacctac	gegeeeeaact	tccacaaaaa	catcaacata
19441	etgettetge	Leeceggeee		gageggaace	acactacat	ccacttcasc
19501	atcctgcaga	gttccctcgg	caacgacctg	egegregacg	gegeeteege	ccgccccgac
19561	agcgtcaacc	tctacgccac	cttcttcccc	arggcgcaca	acaccgcctc	caccctggaa
19621	gccatgctgc	gcaacgacac	caacgaccag	tccttcaacg	actacctctc	ggccgccaac
19681	atgetetace	ccatcccggc	caaggccacc	aacgtgccca	tctccatccc	ctcgcgcaac
19741	tagaccacct	tecacaacta	gagtttcacc	cggctcaaga	ccaaggaaac	tecetecete
19801	ggctcgggtt	togaccccta	ctttgtctac	tcgggctcca	tcccctacct	cgacgggacc
10061	ttctacctca	accacacctt	caagaggtc	tccatcatgt	tcgactcctc	ggtcagctgg
10001	cccggcaacg	accacacacac	caagaaggee	cacttccaca	traagrarag	cattaacaga
19971	eeeggeaacg	accegectect	cacyccyaac	aggeegaga	acttactact	ccadatoctc
19981	gagggctaca	acgrggccca	atgeaacatg	accaaggact	ggttccccgt	ccagacgete
20041	teccactaca	acatcggcta	ccagggcttc	caegtgeeeg	agggetacaa	gyaccycacy
20101	tactccttct	tccgcaactt	ccagcccatg	agcaggcagg	tggtcgatga	gatcaactac
20161	aaggactaca	aggccgtcac	cctacccttc	cagcacaaca	actegggett	caccggctac
20221	cttgcgccca	ccatgcgcca	ggggcagccc	taccccgcca	acttccccta	cccgctcatc
20281	ggctccaccg	cagttccctc	cgtcacccag	aaaaagttcc	tctgcgacag	ggtcatgtgg
20201	cgcatcccat	tetecareaa	ctttatgtcc	atggggggc	tcaccgacct	gggtcagaac
20341	atgctctatg	ccectgoac	ccacacactc	nacatnacct	ttgaggtgga	ccccatggat
20401	gageceacee	testetatet	tatattaas	attttcaaca	tootcagagt	gcaccagccg
20461	gageceaece	tectetatet	cctcttttgaa	geeeegacg	tatecara	caaccctacc
20521	caccgcggcg	teategagge	egiciaceig	cycacycccc	cccccgccgg	caacyccacc
20581	acttaagcat	gagcggctcc	agcgaacaag	agetegegge	categugege	yaccigggat
20641	gcgggcccta	ctttttggga	acccacgaca	agcgcttccc	tggcttcctt	gccggcgaca
20701	agctggcctg	cgccatcgtc	aacacggccg	gccgcgagac	cggaggcgtg	cactggctcg
20761	cctttggctg	gaatccgcgc	tcgcgcacct	gctacatgtt	cgaccccttt	gggttctcgg
20821	accoccooct	caagcagatt	tacagcttcg	agtacgaggc	catgctgcgc	cgaagcgcgc
20881	ttgcctcctc	acccasccac	totctcagcc	tcgagcagtc	cacccagacc	gtgcaggggc
20001	ccgactccgc	cacctacaas	cttttttatt	gcatgtttt	gcatgccttc	gtgcactggc
20941	ccgaccgacc	catacacaca	aaccccacca	traacttroct	gacgggggggg	ссавасооса
21001	tgctacaatc	Catggatgga	atacccacca	taaaaaaaaa	ccaudaudau	ctctaccgct
5100T	tgctacaatc	gccacaggry	tostttssst	ccaygcycaa	ccaggaggag	aaccacage
21121	tcctcgcgcg	ccactcccct	tactttegat	cccaccgcgc	egecategaa	aacgccaccg
21:181	cttttgataa	aatgaaacaa	ctgcgtgtat	ctcaataaac	agcactttat	tttacatgca
21241	ctggagtata	tgcaagttat	ttaaaagtcg	aaggggttct	cgcgctcgtc	gttgtgcgcc
21301	acactagaga	gggccacgtt	geggtaetgg	tacttgggaa	gccacttgaa	ctcggggatc
21361	accagtttgg	gcactggggt	ctcggggaag	gtctcgctcc	acatgcgccg	gctcatctgc
21421	adddcdccca	gcatgtccgg	gccggagatc	ttgaaatcac	aattggggcc	ggtgctctgc
21/21	~222222000	tacaatecec	gagattacan	cactogaaca	ccattagact	ggggtacttc
71E14	acacteres:	. cooggectet	atcactasta	tratecttot	ccaggteete	ggcgttgctc
7 T D 4 T	acaccggcaa	geacyclett	gregeryate	oggeeeeege	adddagaaca	ctraggette
21601	aggccgaacg	gggtcatctt	geacayeegg	cygcccayga	agggcacgcc	ctgaggcttg
21661	tggttacact	cgcagtgcac	gggcarcago	atcatccccg	egeegegetg	catattcggg
21721	tagagggcct	tgacgaaggc	cgtgatctgc	ttgaaagctt	gctgggcctt	agccccctcg
21781	ctgaaaaaca	gaccacact	cttcccgcta	aactoottat	tcccgcaccc	ggcatcatgc
21841	acqcaqcaqc	gcgcgtcatg	gctggtcagt	tgcaccacgo	: tacgtcccca	geggttetgg
21901	gtcaccttge	ccttactaaa	ctgctccttc	aacgcgcgct	gcccgttctc	gctggtcaca
					•	_ <del></del>

21961	tccatctcca	ccacataata	cttgtggatc	atcaccatca	catgcagaca	cttgagctga
22021	ccctcgacat	cacaacaacc	atgatcccac	agggcgcagc	cootocactc	ccagttctta
22021	tgcgcgatcc	cactataact	gaggatgtaa	ccttgcaaca	ggcgacccat	gacggtgcta
22001	aatgctttct	cactactass	gaagacgtac	agaccacaaa	cctcctcatt	catccaggtc
22741	tggcacatct	tttaasaat	ctcaatctac	tcagacataa	gcttgtaagc	atcococago
22201	ccgctgtcga	cccggaagac	ttccatcacc	acattcataa	tatecatece	cttctcccag
22221	gacgagacca	cgcggtagcg	angagatta	cacecattae	dascaccaa	agtcacagg
22321	. gacgagacca . tcgacgatgc	gaggcagact	cagggggttg	ttcaacaca	ccacacacta	actasatece
22381	. tegaegatge . acteceaega	gtttteegte	thateasa	atotottoat	cagaggetg	cttaatcaca
22441	tgcttggtct	ttacggcatc	tteetgggge	accecectet	cggggccac	cacatcatca
22501	tgettggtet	ttetggettg	ccccccccc	tttaaaaaat	taataaaaa	aggeococo
22561	tcggaagacc	eggageecae	cegetgatae	aratagagaga	cggcgggcag	acccaaaac
22621	ggcggcgagg	ggeteetete	etgeteegge	ggatagegeg	tagacccgcg	cattetttee
22681	ggagtggcct	etegetecat	gaaceggege	acgicectgac	aggaggagt	aaccaccac
22741	taggggaaga	tggaggagca	gccgcgraay	ttcaaacac	caactcatct	adaacccca
22801	gagcaaccca	aaatcgagca	ggacctgggc	cccgaagagc	caaccaacac	tagaactccaa
22861	. caggatgaac	aggagcacga	gcaagacgca	gyccaggagg	agaccgacgc	ccaatccatc
22921	catggctacc	tgggaggaga	ggaggatgtg	ctyctaaaac	tcagcatca	agaactatat:
22981	atcctccggg	acgecetgge	cgaccggage	gaaacccccc	ccagegeega	accesecac
23041	cgggcctacg	ageteaacet	ettetegeeg	totagagatat	ttaaacycca	casacacats
23101	. acctgcgagc	ccaacccgcg	teteaaette	nagatagaa	tatestassa	cgaggeceta
23161	gccacctatc	acatetttt	caayaaccaa	aagateeeeg	gastagatga	tatementee
23221	acccgcgccg	acgegeteet	egetetgggg	ataaataaa	acacacaca	caccacces
23283	ctggaagagg	tgeceaagat	cuccyaaggg	coccycleggy	contactact	attanaana
23343	gctctgaaag	aaacagcaga	ggaagagggt	cacactageg	tanagaaatt	cacctacacc
23401	gacaacgcca	ggetggeegt	geteaagege	agegregage	atangetest	catecece
23461	gccgtcaacc	teeegeeeaa	ggteatgegt	cycaccacyy	accageceae	catgeteeac
23523	atcgaggccc	tcgatgaaag	tcaggagcag	egeceegagg	acycccygcc	acaccaccac
23583	. gacgagcagc	tegegegttg	getegggaee	egegaeeeee	aggetttgga	acaycygcyc
23641	aagctcatgc	rggccgrggr	cetggteace	cccyagetcy	aatgeatgeg	cagacacaat
23703	agcgaccccg	agaccctgcg	taaggtegag	gagacectge	actacacter	ctcataccta
23763	ttcgtcaggc	aggcctgcaa	gateteeaae	geggageega	ccaacciggi	creatgeetg
23821	gggatcctgc	acgagaaccg	cctgggacag	accordence	tetecacici	gaagggcgag
23881	gcgcgtcggg	actatgtccg	cgactgtgta	TEECECETE	tengecacac	ccggcaagca
23941	gccatgggcg	tgtggcagca	gtgtetegag	gacyaaaacc	aggagget	ggacaageee
24001	cttgctagaa	accttaaaaa	getgtggaeg	ggerregacg	tananagaa	catacagae
24061	ctggccgaga	tegtttttee	agaacgcccg	aggeagaege	thatagara	atctgggatg
24123	ttcatgagcc	agagcatgtt	gcaaaactac	cycactttca	ccctcgagcg	ccccaactat
24181	ctacccgcca	cctgcaacgc	atteecetee	pactitytee	cyclyayeta	caactacaac
2424	ccccgccgc	tgtggagcca	ctgctatete	ttgeagetgg	ccaactacat	cgcctaccac
2430	tcggacgtga	tcgaggacgt	gagcggcgag	gggetteteg	agreecacre	cegetgeaac
2436.	ctgtgctccc	cgcaccgctc	eetggtetge	aacccccagc	ctctgagcga	gacccaggcc
2442	Lateggtacet	tcgagctgca	aggreegeag	gagtecaceg	ccccgccgaa	acteacycey
2448.	gggttgtgga	cttccgcgta	cctgcgcaaa	tttgtacccg	aggactacca	ctccctcata
2454	ataaagttct	tcgaggacca	ategegeeca	cagcacgegg	ateceaegge	agasttatt
2460	acccagggcg	cgatcctcgc	ccaattgcac	gccatccaaa	aaccccgcca	agagetteett
2466	L ctaaaaaagg	gragaggggr	cracetggae	ceceagaegg	gegaggeget	caacccgggc
2472	L ctcccccagc	atgccgagga	agaagcagga	geegetagtg	gagcagatgg	aayaayaaty
2478	l ggacagccag	gcagaggagg	acgaacggga	ggaggagaca	gaggaggaag	aactyyaaya
2484	l ggtggaagag	gagcaggaaa	cagageagee	egregeegea	ccatccgcgc	eggeageeee
2490	l gccggtcacg	gatacaacct	ccacagetee	ggccaageet	cetegragar	gggategagt
2496	gaagggtgac	ggtaagcacg	ageggeaggg	ctaccgatca	tggagggtee	acaaagecyc
2502:	l gatcatcgcc	tgcttgcaag	actgcggggg	gaacatcgct	tratagecegee	gotacotyct
2508:	l cttccaccgc	ggggtgaaca	tececegeaa	cgtgttgcat	Lactacegic	accucacag
2514	l ctaagaaaaa	gcaagtaaga	ggagtegeeg	gaggaggcct	gaggatcgcg	ycgaacgagc
2520	l cctcgaccac	cagggagctg	aggaaccgga	Letteccae	LCEETATGCC	accet cage
2526	l agagtcgagg	tcagcagcaa	gaactgaaag	caaaaaaccg	gcccccgcgc	cegeecacee
2532	l gcagttgctt	gtaccacaaa	aacgaagatc	agetgeageg	cactetegaa	yacgccgagg
2538	l ctctgttcca	caagtactgc	gegeteacte	ctaaagacta	aggegegeee	acceggaaaa
2544	l aaggcgggaa	ttacctcatc	gecaecatga	gcaaggagat	CCCCACCCCT	acatgtyga
2550	l gctatcagcc	ccagatgggc	cuggeegegg	gegeeteeca	ggactactcc	acceggatga
2556	l actggctcag	rgccggcccc	Legalgalet	cacgggtcaa	eggggteegt	aaccatcgaa

FIG. 16A-7

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05604						
	accagatatt					
25681	gtaattggcc	ctccaccctg	gtgtatcagg	aaatccccgg	gccgactacc	gtactacttc
	cgcgtgacgc					
23801	cttcccggtg	eccyclecge	ccacaacegg	glalaaaaac	cerdara	cyaggcagag
	gcacacagct					
25921	tccaactagc	cggagccggg	agatcgtcct	tcactcccaa	ccaggcctac	ctgaccttgc
	agagcagctc					
	ttgtgccctc					
	ttataccgaa					
26161	gtgactcggc	tgagctcgct	cggttgaggc	atctggacca	ctgccgccgc	ctgcgctgct
	tcgcccggga					
	ctgcacacgg					
	tcacccagca					
	gcatctgtcc					
26461	ataaaagctg	aactaagaac	cttctttgga	atcccttgtc	atcatcaaat	caacaagacc
	atcaacttca					
	atctggtttt					
			_		-	
	ctcctaccta					
	cttcatcgcc					
26761	ttcactttgg	tgaacgttac	cggcagcagc	acagccgctc	cagaaacatc	taaccttctt
	tctgatacta					
	gggagttcta					
	gcagtgctgt					
27001	cattgtgggg	aggaaccatg	aaggggctct	tgctgattat	cctttccctg	gtgggggtg
27061	tgctgtcatg	ccacqaacaq	ccacgatgta	acattaccac	aggcaatgag	aggaacgact
	gctctgtagt					
	ccatgggaaa					
	ctgtccatgg					
	gtgatatcac					
27361	tggtgggttt	ttctttggct	tttgtgatca	tggcctgctt	gatgtcaggt	ctgctggtag
	gggctctagt					
	tgctataaat					
	ctctcttctt					
27601	taatataact	ttagtgggac	cctcagatac	tccagttacc	tggtatgatg	gcaagggatt
27661	gcaattttgt	gacggaagta	cagttaagaa	tccgcagatc	agacatactt	gtaatgatca
	aaacttaact				_	_
	tgacagtaag					
	aaagccacaa					
27901	tggacctcca	ggaattccag	ttagttggta	ttatcataat	ggcacacagt	tctgcgatgg
27961	agataaaatt	attcatccag	aattcaacca	cacctgtgat	aaacaaaacc	ttacactgct
28021	gtttgtaaac	tttacacatq	atggaggeta	tettggatte	aattacaaag	gtactcagag
	aattcagtat					
	_					-
	agaacaaagt					
28201	gggtatagat	acaaatcaaa	agaaagctaa	taacagacaa	aagccatctc	aaaggccatc
28261	aagaagacgg	ccgacaaaca	ctcctgagac	aaaacaactt	acagtgtcta	ttgggtctaa
	cttaacttta					
	atgtgaagaa					
	taatgtaact	_				
	aagatacaga					
28561	cagacctact	actcctgatc	agaaacacag	atttgaatta	caaattgaaa	ataatgcaaa
	tgatgaagaa					
	cttcataact					
	atacaatcat					
28801	ttcagaacca	tgaaggcttt	cacagettge	gttctgttta	acataatcac	acttagtgta
28861	gctgcaaatg	gttttaaaca	tgttaatgtt	accagattaa	gtaatgtaac	actgacagga
	gctggaatta					
	tggatgaata					
	attactaatc					
	gaaagtttta					
29161	attattgagc	ttccaacaac	tagagcaccc	accacagtta	ggacaacaca	gcctaccact
	gtgcccacta					
	J = J = J = D = D D D			_5 5		

FIG. 16A-8

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00001		acacaacagt	gcagaatact	actttattqa	ttgggttttt	actgagagga
29281	acacagetag	ctactgaaca	geagaaaact	acctcaagtg	ccttcagcag	cactgcaaat
29341	aatgaaagta	ttgcttggac	taatgaggee	ggagtatcat	tgatgaatcg	acageettae
29401	ttaacttcgc	atattcaaat	taatgaaacc	attatatata	ggatctttat	tcttgcggtt
29461	tcaggtttgg	atattcaaat	cacciccic	gacgactgtg	ggacacccat	atacaggcca
29521	cttctgtact	ttgtctgctg	caaayccaya	gagaaaccca	taaggaatct	tetettetet
29581	gtaatcgggg	aacctcagcc	tetecaagig	gacggaggee	tatttaecat	cctattctat
29641	tttacagtat	ggtgatcagc	catgattcct	aggittettee	toracac	atatataaaa
29701	ctcttcaaca	tetgtgetge	cttcgcggcc	gtetegeacg	cetegeeeya	cagactaggg
20761	aatttcccaa	catacctcct	ctttacccta	ctaacctgca	cetycytety	Cagcaccgcc
20021	taaataataa	tcacctttct	dcadctcatc	dactddtgct	gegegegeta	Caactacece
20001	caccacacte	cccaatacac	ggacgagaac	gtagccagaa	tettaagget	Cattlyatta
00041		actestacta	atatecetee.	tatecectue	Cultigudact	cccgccgacc
20001	actetaaate	casattegeg	gacatatoga	atttcttaga	ttgctattag	gagaaaaccg
20061	atatacacta.	ctattacttq	gtgattgttg	gggtagtcat	ggtetgetea	Lycactice
20121	++~aaa++a+	matchacece.	tottttaatc	ttaactaaa	Cicigityay	gcartcacac
20101	acacactaca	aaacadttca	ctageeteca	caccaccacc	cacaccycci	ccccycagaa
20041	t	tatrattrar	tacttagaag	adccccctcc	Coggettett	LLCactycta
20201		astsscommo	aacaataact	gaccacctgg	accicgagai	gyacyyccay
30301	gccaccccca	agcgcatcct	gcaactgcgc	gtccgacage	agcaggagcg	ggccgccaag
30301	geeteegage	atgccatcaa	catccaccag	tgcaagaagg	gcatcttctg	cctggtcaag
30421	gageteeteg	tcacctacga	actestates	adcadcaadc	agcatcgcct	cgcctatgag
30481	caggcaaaga	agaagcaaaa	atteacetac	ataataaaca	tcaaccccat	agtcatcacc
30541	ctacccagc	gcgagaccaa	eggetgeete	cacteeteet	acassaaccc	cgagtgcatc
30601	cagcagtcgg	gegagaceaa	thereses	caccacctcc	tecceateaa	ctgatgttga
30661	tactccctcc	tcaagaccct	ttgeggaete	aggasttact	cataacaata	aatcattoga
30721	ttaaaagccc	aaaaaccaat	caaacccttc	cccaattact	catalogates	agtataatta
30781	actaatcatt	caataaagat	cacttacttg	aaatctgaaa	gracycece	ggcgcagccg
30841	ttcagcagca	cctcggaacc	ctcctcccag	ctctggtact	ceagleeceg	gcgggcggcg
20001	anattaataa	acaccttgaa	agggatgtca	aattcctggt	ccacaatttt	Callylllic
20061	antagastag	casagagact	ccaaataaa	gatgacttca	acceegucia	CCCCtatggc
21021	+	atcacaatat	ccccttcctt	actccccct	ttgtttctt	Cyatygatte
21001	cossacttcc	cectagaat	cctotcactc	aaactqqctq	acceaatege	Callactaat
21111		cactcaardt	πααρασσσσ	cttactqttq	adadayatay	tygaaattta
21001		ctaaggctcc	cttocaagtt	acaactgata	aacagttgga	aattycacty
21261	~c+tatccat	ttgaagtcag	taatoocaao	cttggcataa	aaycayytca	Lygattyaaa
21221	~tanttanca	`aaattootoo	tttggaaggt	ttaacaaata	cgcttgtagt	LLLGALLGGA
21201	aaaggaatag	rtactraaaa	tcttgaaaac	agtgatgggt	caagtagagg	ayıtyytata
24441		ttactaaada	tagaggtctg	tettttata	aaaayyytya	Litaging
21501	taasstassa	atratracar	acocactcta	tggacaactc	cegaeceate	CCCaaaccyc
31201	cygaacaaac	aggaaaggga	ttcaaagctc	actttagtat	taacaaaatg	tggcagtcaa
31201	acaaccyacc	atgtctcttt	acttottota	aaaggaaaat	ttagtaacat	aaacaataat
31621	attttggtta	ctgataaaaa	actogoogta	aagctacttt	ttaatgaaaa	gggagtatta
31681	actaatccaa	cgacacttaa	gaccacagea	togaactaca	gaaatgataa	ttctactqta
31741	arggacagu	atgataatgc	antegatet	atoccasaca	taaaagetta	tcctaaacct
31801	tctcaggcct	cttcggctaa	ageceeeee	acgeedaded	ctactaaaaa	atacattoto
31861	accacagaca	cttcggctaa	accayaayac	aaaaaaaagtg	ttataactat	taagtttaat
31921	agcaatgtct	atattggagg	CLigidayac	tttaatta	cataaacaaa	aacctttgaa
31981	gcagaaactg	aatgtgctta	ttcgattacc	ttttaatata	ttacccaaca	asstractac
32041	gatgtgcagt	ttgattcctc	etetttaee	tetatata	tottaatt	tacaccacca
32101	gaagacaaat	aaaatgttt	aaaatgaatt	catguatett	carryattet	atcaccagea
32161	. cgggtagtca	gtctcccacc	accagcccat	ttcacagtgt	aaacgattet	ctcagtatgg
32221	gtggccttaa	atagggaaat	gttctgatta	gtgcgggaac	tggacttggg	gictataatc
22201	cacacacttt	cctaacaaac	caaacddddd	tcqqtgattg	agatgaagco	giccicigaa
22211	aagtcatcca	accodocte	acagtccaag	gtcacagtct	ggtgaaacya	. yaayaacyca
22401	angettcata	ctcccaaaaac	aggatgggtC	tataccicio	carcagegee	Cicaacagic
22461	+ataccacca	, aaactcaata	caactactac	agatgggatc	gggatcacaa	giciciciga
20521	atataataa	, cacagootto	agcatcagto	tcctggtgcg	tegggeacag	cacegeatee
22501	testatagat	· catottetea	cagtaagtgg	: agcacataat	: caccatgtta	tteageagee
22641	antestra:	, aatactccaa	ccaaaactca	l tottggggat	. gatggaaccc	acytyaccat
22701	- cetaccadat	- acaacaatat	atcagatgcc	tocccccat	gaacacactg	CCCatataca
22761		- gaggatatet	ctottcacaa	l tctgacggta	l ccagggaaag	, cgctggttga
22021	agatgcacco	r gtaaatgact	ctectgaacc	: acacggccac	cagggtgcct	. eccycecyac
32021	acatycaeco	gcccggggat	gaacagtgg	aatgcaggat	ccagcactca	taccegetca
27887	Lactycayyy	. gcccygygat	. 3~~~~			•

FIG. 16A-9

32941	ccatctgagc	tctcaccaag	tccagggtag	cggggcacag	gcacactgac	atacatcttt
33001	ttaaaatttt	tatttcctct	ggagtcaaga	tcatatccca	ggggactgga	aactcttgga
33061	gcagggtaaa	gccagcagca	catggtaatc	cacggacaga	acttacatta	tgataatctg
33121	catgatcaca	atcaggcaac	aggggatgtt	gttcagtcag	tgaagccctg	gtttcctcat
33181	cagatcgtgg	taaacqqqcc	ctgcgatatg	gatgatggcg	gagcgagctg	gattgaatct
33241	cggtttgcat	tgtagtggat	tctcttgcgt	accttgtcgt	acttctgcca	gcagaaatgg
33301	gcccttgaac	agcagatacc	cctcctgcgg	ccgtcctttc	gctgctgccg	ctcagtcatc
33361	caactgaagt	acatccattc	tcgaagattc	tggagaagtt	cctctgcatc	tgatgaaaca
33421	aaaaacccgt	ccatgcgaat	tcccctcatc	acatcagcca	ggactctgta	ggccatcccc
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33541	atttttattc	caaacggtct	cgaaggacga	taaagtgcaa	gtcacgcagg	tgacagcgtt
33601	cccctccgct	gtgctggtgg	aaacagacag	ccaggtcaaa	acccactcta	ttttcaaggt
33,661	gctcgaccgt	ggcttcgagc	agtggctcta	cgcgtacatc	cagcataaga	atcacattaa
33721	aggctggccc	tccatcgatt	tcatcaatca	tcaggttaca	ttcctgcacc	atccccaggt
33781	aattctcatt	tttccagcct	tggattatct	ctacaaattg	ttggtgtaag	tccactccgc
33841	acatotogaa	aagctcccac	agtgccccct	ccactttcat	aatcaggcag	accttcataa
33901	tagaaacaga	tcctgctgct	ccaccacctg	cagcgtgttc	aaaacaacaa	gattcaataa
33961	aattctaccc	tccgccctga	gctcgcgcct	caatgtcagc	tgcaaaaaat	cacttaagtc
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34081	ggaaaacttt	aatgctccaa	agctagcacc	caaaaactgc	atgctggaat	aagctctctt
34141	tgtgtctccg	gtgatgcctt	ccaaaatgtg	agtgataaag	cgtggtagtt	tttctttaat
34201	catttgcgta	atagaaaagt	cctgtaaata	agtcactagg	accccaggga	ccacaatgtg
34261	gtagcttaca	ccgcgtcgct	gaagcatggt	tagtagagat	gagagtctga	aaaacagaaa
34321	gcatgcacta	aactaaggtg	gctattttca	ctgaaggaaa	aatcactctc	tccaacaaca
34381	gggtacccac	tgggtggccc	ttgcggacat	acaaaaatcg	gtccgtgtga	ttaaaaagca
34441	gcacagtaag	ttcctgtctt	cttccggcaa	aaatcacatc	ggactgggtt	agtatgtccc
34501	tggcatggta	gtcattcaag	gccataaatc	tgccctgata	tccagtagga	accagcacac
34561	tcacttttag	gtgaagcaat	accaccccat	gcggaggaat	gtggaaagat	tcagggcaaa
34621	aaaaattata	tctattgcta	gtcccttcct	ggacgggagc	aatccctcca	ggactatcta
34681	tgaaagcata	cagagattca	gccatagctc	agcccgctta	ccagtagaca	gagagcacag
34741	cagtacaagc	gccaacagca	gcgactgact	acccactgac	ccagctccct	atttaaaggc
34801	occttacact	gacgtaatga	ccaaaggtct	aaaaaccccg	ccaaaaaaaa	acacacacgc
34861	cctgggtgtt	ttttgcgaaa	acacttccgc	gttctcactt	cctcgtattg	atttcgtgac
34921	ttaacttccg	ggttcccacg	ttacgtcact	tctgccctta	catgtaactc	agtcgtaggg
34981	cgccatcttg	cccacgtcca	aaatggcttc	catgtccagc	cacgcctccg	cggcgaccgt
35041	tagccgtgcg	tcgtgacgtc	atttgcatca	tcttctctcg	tccaatcagc	gctggccccg
35101	ccctaaattc	aaaagctcat	ttgcatgtta	acttttgttt	actttgtggg	gtatattatt
35161	gatgatc					
SEQ I	D NO: 5		•			

Grp	Vaccine	Monkey	Р	re	W	k 4	W	k 8	W	12_
•	at Wk 0, Wk 4	. ID	Mock	Gag	Mock	Gag	Mock	Gag	Mock	Gag
1	Ad24AE 1gcgAOrf8Ad5Orf6	00C072	3	4	4	381	3	150	3	68
	10^11 vp	00C178	3	3	1	559	1	743	0	635
	•	00C222	0	3	1	369	1 1	753	0	670
		00D011	1	9	9	211	4	273	0	520
•		00D023	0	6	0	295	1	459	1	368
		00D031	15	5	10	103	1	101	1	40
2	Ad24AE 1gogAOrf6Ad5Orf6	99C168	4	6	0	118	5	241	3	209
	10^10 vp	99C170	10	5	5	241	3	141	3	103
		99C173	1	3	°	23	٥	14	0	21
3	Ad24AE 1gogAE4Ad5Orf6	99C154	0	3	0	93	0	60	1	53
	10^10 vp	99C158	1	0	1	141	0	101	1	120
		99C177	0	0	٥	45	0	39	0	79
4	MRKAd5-HIVgag	00C018	1	5	13	1025	0	824	3	753
	10^11 vp	00C034	0	4	5	219	5	404	0	491
		00C058	4	4	3	1086	0	440	0	439
5	MRKAd5-HIVgag	99C218	0	3	5	2500	0	1580	10	1655
•	10^10 vp	99C227	6	1	4	529	5	365	5	1004
		99D185	ND	ND	0	425	0	310	0	271

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Vaccine	Monkey	Gag-Specific (Wk 12)		
at Wk 0, Wk 4	ID_	%CD4	%CD8	
Ad24AE 1 gagAOrf6Ad5Orf6	00C072	0.02	0.02	
10^11 vp	00C178	0.05	0.38	
	00C222	0.02	0.40	
·	00D011	0.02	0.27	
	00D023	0.01	0.11	
	00D031	0.01	0.01	
MRKAd5-HIVgag	00C018	0.05	0.41	
10^11 vp	00C034	0.06	0.18	
	00C058	0.02	0.28	

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Grp	Vaccine	Monkey	Wk 4	WK 8
	at Wk 0, Wk 4	ID		
1	Ad24ΔE 1 gagΔOrf6Ad5Orf6	00C072	<10	77
	10^11 vp	00C178	<10	26
	•	00C222	<10	423
		00D011	<10	98
		00D023	<10	<10
		00D031	<10	<10
2	Ad24AE 1 gogAOrf6Ad5Orf6	99C168	<10	<10
<b>,</b>	10^10 vp	99C170	<10	<10
	•	99C173	<10	<10
	A 104453 AF44450-10	200454	10	
3	Ad24∆E1gcg∆E4Ad5Orf6	99C154	<10	<10
	10^10 vp	99C158	<10	<10
•		99C177	<10	<10
4	MRKAd5-HIVgag	00C018	34	1017
	10^11 vp	00C034	14	423
,		00C058	46	934
5	MRKAd5-HIVgag	99C218	20	99
	10^10 vp	99C227	40	767
		99D185	17	342

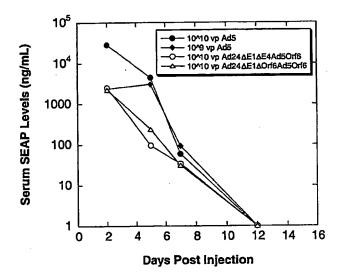


FIG. 20

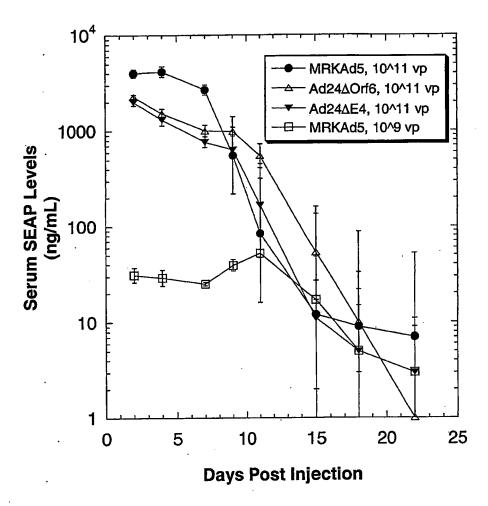
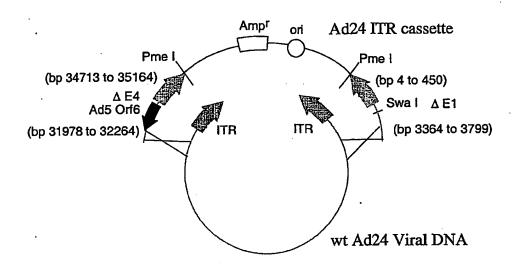


FIG. 21

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Animai	Prime (Wk 0, 4, 26)	(k 0. 4. 26) Boost (Wk 56)		Pre		Prime <sup>b</sup>		Pre-Boost <sup>e</sup>		Post-Boost	
Ammai.	t time (ask of al no)		Mock*	Gag*	Mock	Gag	Mock	Gag	Mock	Gag	
Monkey 1	10° vp MRKAd5-gag	1011 vp Ad24ΔE1gagΔOrf6Ad5Orf6	18	16	1	244	3	74	3	1235	
Monkey 2	107 vp MRKAd5-gag	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf8Ad5Orf8	10	9	4	83	0	18	٥	856	
Monkey 3	10° vp MRKAd8-gag	1011 vp Ad24ΔE1gagΔOrf6Ad5Orf6	l 1	1	0	219	9	69	0	703	
Monkey 4	10 <sup>7</sup> vp MRKAd8-gag	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6	1	1	3	59	1	20	0	419	
Mankey 5	none	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf8	3	4	ND*	ND	ND	ND	4	558	
Monkey 6	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	٥	3	ND	ND	ND	ND	1	295	
Monkey 7	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	1	9	ND	ND	ND	ND	8	103	
Monkey 8	none	1011 vp Ad24AE1gagAOr/6Ad5Or/6	3	3	ND	ND	ND	ND	1	381	
Monkey 9	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	0	6	ND	ND	ND	ND	0	369	
Monkey 10	none	1011 vp Ad24AE1gagAOrf6Ad5Orf6	15	5	ND	ND	ND	ND_	10	211	

Animal	Prime (Wk 0, 4, 26)	Boost (Wk 56)	Gag-Specific T cells (Wk 60)		
Aillinai	t tano (tra o) + 1 20)		%CD4	%CD8	
Monkey 1	109 vp MRKAd5-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.06	0.37	
Monkey 2	107 vp MRKAd5-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.01	0.56	
Monkey 3	10 <sup>9</sup> vp MRKAd6-gag	10 <sup>11</sup> vp Ad24ΔE1gagΔOrf6Ad5Orf6	0.07	0.06	
Monkey 4	10 <sup>7</sup> vp MRKAd6-gag	10 <sup>11</sup> vp Ad24∆E1gag∆Orf6Ad5Orf6	0.04	0.20	

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Animal	Prime (Wk 0, 4)	Boost (Wk 24)	Pre		Prime <sup>b</sup>		Pre-Boost <sup>e</sup>		Post-Boost <sup>d</sup>	
	, , ,	·	Mock	Gag*	Mock	Gag	Mock	Gag	Mock	Gag
Monkey 11	1011 vp Ad24AE1gagAOrf6Ad5Orf6	107 vp MRKAd5-gag	3	4	3	150	4	28	0	188
Monkey 12	1011 vp Ad24AE1gagAOrf8Ad5Orf6	107 vp MRKAd5-gag	0	3	1	753	4	554	0	1029
Monkey 13	1011 vp Ad24ΔE1gagΔOrl6Ad5Orl6	10 <sup>7</sup> vp MRKAd5-gag	1	9	4	273	0	370	0	1520
Monkey 14	none	10 <sup>7</sup> vp MRKAd5-gag	0	0	ND*	ND	ND	ND	4	94
Monkey 15	none	107 vp MRKAd5-gag	0	0	ND .	ND	ND	ND	1	168
Monkey 16	enon	107 vp MRKAd5-gag	8	3	ND	ND	_ND	ND_	_ 8	149

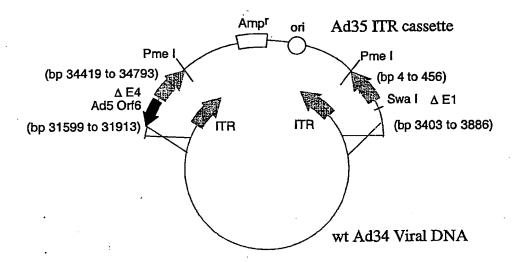
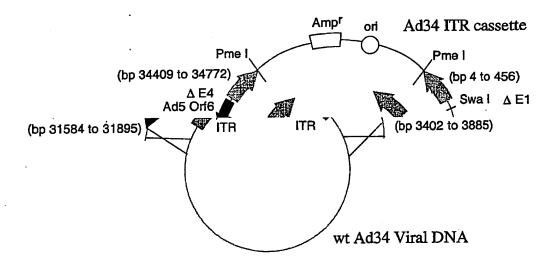


FIG. 26



_	_					
1	catcatcaat	aatatacctt	atagatggaa	tggtgccaat	atgtaaatga	ggtgatttta
61	aaaattotoo	aatatataat	gattggctgt	aganttaaca	actasacaaa	acaacacaac
121	aataaaaaa	550505055				909909090
121	cycgggaaaa	Lgacgillig	tgggggtgga	gttttttgc	aagttgtcgc	gggaaatgtg
181	acgcataaaa	aggctttttt	tctcacggaa	ctactgactt	ttcccacggt	atttaacagg
.241	aaatgaggta	gttttgaccg	gatgcaagtg	aaaattocto	atttgcgcgc	gaaaactgaa
301	taeaasaata	tttttctcaa	taatgtggta	tttatagoog	actesostat	ttattanaaa
301	rgaggaagrg	LLLLLLLyaa	caacycygra	LLLauggeag	ggrygagrar	Ligiteaggg
3 P T	ccaggtagac	tttgacccat	tacgtggagg	tttcgattac	cgtgtttttt	acctgaattt
421	ccacatacca	totcaaaotc	ttctgttttt	acotacotot	cagctgatcg	ctacootatt
481	tatacctcag	aatttatata	aagaggccac	tattasataa	30-3-0-3	actttctcc
E 41	tataccccag	ggcccgcc	aagaggccac	ccccgagtgc	cagcyayaay	agriculture
241	rergegeegg	cagtttaata	ataaaaaaat	gagagatttg	cgatttctgc	ctcaggaaat
601	aatttctgct	gagactggaa	atgaaatact	ggagcttgtg	gtgcacgccc	tgatgggaga
661	cgatccggag	ccacctgtgc	agctttttga	acctectaca	cttcaggaac	totatoattt
721	agagg agag	acatocaaca	attataataa	goodcotatg	ceteaggaae	thearabha
721	ayayytayay	gyarcygagg	attctaatga	ggaagergrg	aatggctttt	ttaccgattc
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			ctgcagcggg			
			ttcctggaca			
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1141	tatttacagt	aagtgtgttt	aagttaaaat	ttaaannaat	atactatttt	tracatotat
1201	attenetes	225	thethethet		t-t-t-t-	ccacatgtat
1201	arryagrygg	agttttgtge	ttcttattat	aggreergrg	tetgatgetg	atgagtcacc
1261	atctcctgat	tctactacct	cacctcctga	gattcaagca	cctgttcctg	tggacgtgcg
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1.001	5-5	bbasses	agaaayacta	ggcaactgtt	ayayyacycc	ccggacggag
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1861	csaccccauu	tagaactgcc	gctgctgtgg	cttttcttcc	+++++++	antontan
1001	taaccccagg	tagaactgcc	gergergeg	Cultural	LLLLacatta	yacaaacyya
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23/11	antttctnta	ttageageage	aatattcact	gazzgaggta	222222	gaaggaaga
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24UI	tgaggatgat	tgggaggtgg	ccattaaaaa	ttatgccaag	atagctttga	ggcctgataa
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2521	gactgagata	gtaatagata	ctcaagacaa	ggcagttatt	agatoctoca	tgatggatat
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2201	geggeeegga	gragicygra	tggaagcagt	aactitigia	aatyttaayt	rraggggaga
2641	tggttataat	ggaatagtgt	ttatggccaa	taccaaactt	atattgcatg	gttgtagctt
2701	ttttggtttc	aacaatacct	gtgtagatgc	ctggggacag	gttagtgtac	ggggatgtag
2761	tttctatgcg	tattagatta	ccacagctgg	cadaaccaad	antcaattot	ctctcaacaa
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3181	cagaatgagc	ctaacaggaa	tctttgacat	gaacatgcaa	atctggaaga	tcctgaggta
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3301	-2240409	ataastatas	atananti-	-gaacycyga	ast the entire	the
22C1	googgegege	yrayargtga	ctgaagatct	yayaccggat	catttggtta	LIGCCCGCAC
3361	rggagcagag	ttcggatcca	gtggagaaga	aactgactaa	ggtgagtatt	gggaaaactt
3421	ggggtggggt	tttcagatgg	acagattgag	taaaaattto	ttttttctct	ctttcagctg
3481	tcatgagtgg	agacocttct	tttaaggggg	gagtettese	cccttatatata	acadecate
35/11	tocostast -	~~~~~	actoococt	beatering		acayyycytc
JJ41	LUCCATCCTG	ygcaggagtt	cgtcagaatg	ttatgggatc	tactgtggat	ggaagacccg
3601	TCCaacccgc	caattcttca	acgctgacct	atgctacttt	aagttcttca	cctttggacg
3661	cagctgcagc	caccaccacc	gcctctgttg	cccctaacac	tatacttaca	atgggttact
3721	atquantat	cataactaat	tccacttcct	ctaataacco	ttctaccet~	
3701	arttactte	antiti	asaataas			acceaggaca
2041	ayccacttgt	CCLELEGGCC	cagctggagg	ctttgaccca	acgtctgggt	gaactttatc
<b>3841</b>	agcaggtggc	cgagttgcga	gtacaaactg	agtctgctgt	cggcacggca	aagtctaaat
				•	•	_

FIG. 28A-1

	_					
3901	aaaaaaaaat	tccacaatca	atgaataaat	aaacgagctt	gccgccgacc	taaaatcaay
3961	tgtttttatt	tcatttttcg	cgcacggtat	gccctagacc	accgatctcg	atcattgaga
4001	acacggtgga	ttttttaaa	astrotatao	aggragatt	gaatgtttag	atacatgggc
4021	acacggugga	ttttttttag	auccecucug	4990999400	attactacta	cacatagta
4081	attaggccat	ctttggggtg	gagatagete	cattgaaggg	atteatgete	cggggtagtg
4141	ttgtaaatca	cccagtcata	acaaggtcgc	agtgcatggt	gttgcacaat	atcttttaga
4201	agtaggctga	ttaccacaga	taagcccttg	gtgtaggtgt	ttacaaacco	attaaactaa
42 U I	aytayyctya	t tt		2030-33030	ccatttttaa	attaccasts
4261	gaggggtgca	ttcggggtga	aaccargrac	accinggact	ggattttaa	gitggcaata
4321	ttgccgccaa	gatctcgtct	tgggttcatg	ttatgaagga	ccaccaagac	ggtgtatccg
1381	gtacatttag	gaaatttatc	atataactta	gatggaaaag	cataaaaaa	tttggagaca
4301	gracartrag		ttaataaa	testeestes	taatammaat	aggaccataa
444 <b>1</b>	cccttgtgtc	ccccgagact	Liccarycac	Caccacga	caacagcaac	ggggccgcgg
4501	gcagcagcgc	gggcaaacac	gttccgtggg	tctgacacat	catagttatg	tteetgagtt
4561	aaatcatcat	aagccatttt	aatgaatttg	ggggggggg	tacccgattg	gggtatgaat
4601	gttccttcgg	addecadade	atacttcccc	tcacagattt	gcatttccca	agettteagt
4621	greeerregg	geeeeggage	acageeeeee			+
4681	tccgatggtg	gaatcatgtc	cacciggggg	gctatgaaga	acaccyttte	rggggcgggg
4741	gtgattagtt	gggatgatag.	caagtttctg	agcaattgag	atttgccaca	tccggtgggg
4801	ccataaatga	ttccgattac	aggttgcagg	togtagttta	gggaacggca	actgccgtct
4061	tctcgaagca	244444444	ctcattcatc	atttccctta	catgcatatt	ttcccgcacc
4001	tetegaagea	aggggccac	tbb-		*****	ccccgcact
4921	aaatccatta	ggaggcgctc	teeteetagt	gatagaagtt	Citylagiga	yyaaaayttt
4981	ttcagcggtt	ttagaccgtc	agccatgggc	attttggaga	gagtttgctg	caaaagttct
5041	agtctgttcc	acagttcagt	gatgtgttct	atggcatctc	gatccagcag	acctcctcgt
5041	ttcgcgggtt	teeseeste	atagagtaaa	atetaereca	atagacatec	adcactacca
PIOT	ttegegggtt	Lggacggctc	cuyyay cayy.	gcatgagacg	t-t-t-	agegeegeea
5161	gggttcggtc	cttccagggt	ctcagtgttc	gagtcagggt	tgtttccgtc	acagtgaagg
5221	gatatacacc	tacttagaca	cttqccaqqq	tgcgcttcag	actcattctg	ctggtggaga
5201	acttctgtcg	attagagaga	tatatataa	ccaagtagca	gtttaccatg	agttcgtagt
2201	acticiging	Cutggugue		ccaagtagca	tttaaaaatt	ttattaata
5341	tgagcgcctc	ggctgcgtgg	cctttggcgc	ggagerrace	LLLygaagit	Lucitycata
5401	ccgggcagta	taggcatttc	agcgcataca	gcttgggcgc	aaggaaaatg	gattctgggg
5461	agtatgcatc	tacaccacaa	gaggcgcaaa	cagtttcaca	ttccaccagc	caggttaaat
5501	ccggttcatt	aaaataaaa	acaacttttc	coccatattt	tttgatgcgt	ttcttacctt
2277	eeggtteatt	ggggttaaaa	acaageeeee	tereseases		tececators
5581	tggtctccat	gagttcgtgt	cctcgttgag	tgacaaacag	getgteegta	cccccgcaga
5641	ctgattttac	aggcctcttc	tccagtggag	tgcctcggtc	ttcttcgtac	aggaactctg
5701	accactctga	tacaaaaaca	cacatecaga	ccagcacaaa	ggaggctatg	taggaggggt
5701	agcgatcgtt		~~~taa	tttocaaaact	250000000	atateaceet
2/61	agcgatcgtt	gccaaccagg	gggtccacct	LLLCCaaayL	atycadacac	acyccacccc
5821	cttcaacatc	caggaatgtg	attggcttgt	aggtgtattt	cacgtgacct	ggggtccccg
5881	ctagagagat	ataaaagggg	geggttettt	gctcttcctc	actgtcttcc	ggatcgctgt
50/1	ccaggaacgt	carctattaa	ggtaggtatt	ccctctcgaa	aacaaacata	acctctgcac
3341	ccaygaacgc	cageegeegg	9900990000	atttattat	ggogggcccg	attaeastac
6001	tcaggttgtc	agtttctaag	aacgaggagg	alligatati	gacagraceg	guigagaigu
6061	ctttcatgag	gttttcgtcc	atttggtcag	aaaacacaat	ttttttattg	tcaagtttgg
6121	tggcaaatga	tccatacagg	gcgttggata	aaagtttggc	aatggatcgc	atggtttggt
6101	tetttteett	atconcacac	tetttageag	coatottoao	ttogacatac	tegegtgeta
0101		greegege	-tthetas	character	anagattata	acttacasa
6241	ggcacttcca	ttcggggaag	atagttgtca	atteatetgg	cacgattete	actigodace
6301	ctcgattatg	caaggtaatt	aaatccacac	tggtggccac	ctcgcctcga	aggggttcgt
6361	tggtccaaca	gaggetaget	cctttcctag	aacagaaagg	aggaagtagg	tctagcataa
6421	gttcatcggg	agetetee	tocatootaa	agattcccgg	aagtaaatcc	ttatcasast.
0421	grecareggy	agggtttgta	tetatggeaa	agacccccgg	tageautee	
6481	agctgatggg	agtggggtca	tctaaggcca	tttgccattc	tegagetgee	agracacact
6541	catatgggtt	aaggggactg	ccccagggca	tgggatgggt	gagtgcagag	gcatacatgc
6601	cacagatgtc	atagacgtag	atgggatect	caaagatgcc	tatataggtt	ggatagcatc
6661	gccccctct		accacatact	catatecttc	atataataac	actagrasco
000T	geeeecetet	yatactiget	the metatage	-b-bb-b-b-	~~~	5000500000
6721	ccggacccaa	gttggtgcga	regggerere	ccgtcctgta	yacaatctyg	cyaaayacyg
6781	cgtgagaatt	ggaagagatg	gtgggtcttt	gaaaaatgtt	gaaatgggca	tgaggtagac
6841	ctacagagtc	teteacaaag	toggcataag	attettgaag	cttggttacc	agttcggcgg
6001	tgacaagtac		cogtactcaa	atatttatta	aatratra	taacctoott
990T	Lyacaagtac	gcccagggcg	Lagragecaa	gracere	therestee	theeset
6961	ggtttttctt	ttcccacagt	tegeggttga	gaaggtatte	ttegegatee	LLCCaglact
7021	cttctagcgg	aaacccgtct	ttgtctgcac	ggtaagatcc	tagcatgtag	aactgattaa
7081	ctgccttgta	adddcadcad	cccttctcta	caaataaaaa	gtatgcttga	gcagcttttc
7111		~5~~~5~~~	accasactat	atatasaast	gactttgaga	aattootatt
1141	gcagcgaagc	gradiaagg	gcgaaggcgt	cicigacial	yacttuyaya	autoggiati
7201	tgaagtccat	gtcgtcacag	gctccctgtt	cccagagttg	gaagtctacc	cgtttcttgt
7261	aggcggggtt	gggcaaagcg	aaagtaacat	cgttgaagag	aatcttaccg	gctctgggca
7221	taaaattgcg	agtgatggg	aaaggctgtg	gtacttcccc	tegattotto	atcacctoon
1267	t-	ag cyatytyy	anaggeogeg	5-4-6-6-6-6-6-6-6-6-6-6-6-6-6-6-6-6-6-6-	+	2244040499
7381	cagctaggac	gatetegteg	aaaccgttga	tgttgtgtcc	Lacgatgtat	aattetatga
7441	aacgcggcgt	gcctttgacg	tgaggtagct	tattgagctc	atcaaaggtt	aggtctgtag
7501	ggtcagataa	gacateatat	tegagageee	attcotocao	gtgaggattt	gcatgtagga
	atgatgacca					
120T	acyacyacca	aayacccacc	goodg tycty	cuyuaauug	guccuyatat	tattatta
7621	gctggccaat	tgccattttt	tetggagtga	cacagtagaa	ggrrcrgggg	rettgttgcc
7681	atcgatccca	ctttagttta	atggctagat	cgtgggccat	gttgacgaga	cgctcttctc
	_	. •				
7741	ctgagagttt	catgaccago	acgaaaggaa	CTACTTULL	gccaaaqqac	cccatccadd

FIG. 28A-2

7801	tgtaagtttc	cacategtag	gtcaggaaga	atctttctat	gcgaggatga	gageegateg
7861	ggaagaactg	gatttcctgc	caccagttgg	aggattggct	gttgatgtga	tggaagtaga
7921	agtttctgcg	acacaccasa	cattcgtgtt	totocttota	cagacggccg	cagtagtcgc
7021	agcgttgcac	agattatata	tcgtgaatga	actataccta	acttccctta	acgagaaatt
9041	tcagtgggaa	accasacct	accattata	teteatacte	ttctatattc	actatateaa
0101	cctgttcatc	ttatattta	ataataataa	tactaacaaa	ccccacaa	aggcaagtcc
0101	agacctcggc	accasasas	graggrage	casccadad	acacegaggg	gagetgteea
8101	agaeetegge	gegggagggg	cygayctyaa	tagatagage	caraaratta	acttocatoa
8221	gagtcctgag	acgctgcgga	cicagginag	cayytayyya	ttccacacct	teatttataa
828I	tcttttccag	ggcgcgcggg	agguedayau	ggtacttgat	coccacagge	cotttattt
8341	agatgtcaat	ggcttgcagg	gttccgtgtc	ettegggege	gagaaggaat	cccccgccc
8401	ttcttttgat	cggtggtggc	tetettgett	erigeatget	cayaaycyac	gacggggacg
8461	cgcgccgggc	ggaagcggtt	gttccggacc	eggaggearg	gerggrageg	geacgupgge
8521	gccgcgcacg	ggcaggttct	ggtactgcgc	tetgagaaga	ettgegtgeg	teaccacgeg
8581	tcgattgacg	tcttgtatct	gacgtctctg	ggtgaaagct	accggccccg	tgagettgaa
8641	cctgaaagag	agttcaacag	aatcaatttc	ggtatcgtta	acggcagctt	gccccagtat
8701	ttcttgtacg	tcaccagagt	tgtcctggta	ggcgatctcc	gccatgaact	getegatete
8761	ttcctcctga	agatctccgc	gacccgctct	ctcgacggtg	gccgcgaggt	cattggagat
8821	acggcccatg	agttgggaga	atgcagtcat	geeegeeteg	ttccagacgc	ggctgtaaac
8881	cacggccccc	tcggagtctc	ttgcgcgcat	caccacctga	gcgaggttaa	gctccacgtg
8941	tctggtgaag	accgcatagt	tgcataggcg	ctgaaaaagg	tagttgagtg	tggtggcaat
9001	gtgttcggcg	acgaagaaat	acatgatcca	tcgtctcagc	ggcatttcgc	tgacatcgcc
9061	cagagettee	aagcgctcca	tggcctcgta	gaagtccacg	gcaaaattaa	aaaactggga
9121	atttcacaca	gacacggtca	attcctcctc	gagaagacgg	atgagttcgg	ctatggtggc
9181	ccgtacttcg	cgttcgaagg	ctcccgggat	ctcttcttcc	tcttctatct	cttcttccac
9241	taacatctct	tcttcgtctt	caggcggggg	cggagggggc	acacggcgac	gtcgacggcg
9301	cacgggcaaa	cggtcgatga	atcgttcaat	gacctctccg	cggcggcggc	gcatggtttc
9361	agtgacggcg	caaccattct	cgcgcggtcg	cagagtaaaa	acaccgccgc	gcatctcctt
9421	aaagtggtga	ctgggaggtt	ctccgtttgg	gagggagagg	gcgctgatta	tacattttat
9481	taattggccc	gtagggactg	cgcgcagaga	tctgatcgtg	tcaagatcca	cgggatctga
9541	aaacctttcg	acgaaagcgt	ctaaccagtc	acagtcacaa	ggtaggctga	gtacggcttc
9601	ttgtgggcgg	gggtggttat	atattcaatc	tagatettet	gtttcttctt	catctcggga
9661	aggtgagacg	atactactaa	toatoaaatt	aaagtaggca	gttctaagac	ggcggatggt
9721	ggcgaggagc	accapatett	tagatccaac	ttoctogata	cocaoocoat	tggccattcc
9791	ccaagcatta	tectageate	taggaagatc	tttgtagtag	tettecatea	accattctac
00/1	gggcacttct	tecteaceea	ttctaccata	catacgtgtg	agtccaaacc	cococattoo
0001	ttgtaccagt	acceaatcea	ctacgactct	ttcaacaaaa	atggcttgct	gtacttgggt
0061	gagggtggct	trasartrat	caaaatccac	aaaggggtgg	taagccccgg	tattaatggt
10021	gtaagcacag	ttaaccataa	ctgaccagtt	aactotctoo	tgaccaggg	gcacgagete
10021	ggtgtattta	aggccacga	aggeouget	atcaaagatg	taatcottoc	aggtgcgcac
10101	cagatactgg	taacctataa	aggegeggge	caataattaa	caatagaaga	gccatcgttc
10201	tgtagctgga	anagagaga	casaatette	caacataadd	cantastaac	catagatata
T050T	cctggacatc	gegeeggggg	chaggicit	antanaance	caaaaaact	cacatacaca
T070T	cetggacate	theetee	gastgasgata	agtagaagte	agaggaaact	raccartrar
T035T	gttccaaatg	ttgcgtageg	gcacgaagca	grecategea	ggcacggttt	accagingag
10381	gcgcgcgcag	tcattgatge	tetatagaca	cygagaaaat	gaaagegee	tagagaatta
10441	ctccgtagcc	tggaggaacg	tgaacgggtt	gggtegeggt	gtaccccggt	coagactty
10501	tactcgagcc	ggccggagcc	geggetaacg	tggtattggt	teeeestee	cgacccagcc
10561	tacaaaaatc	caggatacgg	aatcgagtcg	ttttgetggt	tgeegaatgg	tagggaagtg
10621	agtcctattt	tttttttt	ccgctcagat	gcatcccgtg	ctgcgacaga	tgegteecea
10681	acaacagccc	ccctcgcagc	agcagcaacc	acaaaaggct	gtccctgcaa	ctactgcaac
10741	tgccgctgtg	agcggtgcgg	gacagecege	ctatgatctg	gacttggaag	agggcgaagg
10801	actggcacgt	ctaggtgcgc	cttcgcccga	gcggcatccg	cgagttcaac	tgaaaaaaga
10861	ttctcgcgag	gcgtatgtgc	cccaacagaa	cctatttaga	gacagaagcg	gcgaggagcc
10921	ggaggagatg	cgagcttccc	gctttaacgc	gggtcgtgag	ctgcgtcacg	gtttggacag
10981	aagacgagtg	ttgcgggacg	aggatttcga	agttgatgaa	gtgacaggga	tcagtcctgc
11041	caggggagagag	ataactacaa	ccaaccttgt	atcggcttac	gaacagacag	taaaggaaga
11101	gcgtaatttc	caaaagtctt	ttaataatca	tgtgcgaacc	ctcattgccc	gcgaagaagt
11161	caccetteet	ttgatgcatt	tataggattt	gatggaagct	atcattcaga	accctactag
11221	caaacctctg	accocacaoc	tatttctaat	ggtgcaacac	agcagagaca	atgaggcttt
11281	cagagagggg	ctgctcaaca	tcaccgaacc	cgaggggaga	tggttgtatg	atcttatcaa
11341	cattetacag	agtatcatag	tgcaggagcg	gagcctgggc	ctggccgaga	aggtggctgc
11401	catcaattac	tcaattttaa	gcttgggaaa	. gtattacgct	cgcaagatct	acaagactcc
11461	atacetteee	atagacaagg	aggtgaagat	agatgggttc	tacatacaca	tgacgctgaa
11501	agtattaeca	ctgagcgatg	atcttoccot	gtaccgcaat	gacagaatoc	atcgcgcggt
11581	gagcgccacc	addadacaca	agttaagcga	cagggaacto	atgcacagtt	tgcaaagagc
11641	totasotore	actagaacca	agggtgagaa	ttactttgat	atgggagctg	acttgcagtg
TT02T	Jetuaetyga	200234466	-555050500			

11701	gcagcctagt	cgcagggctc	tgaacgccgc	gacggcagga	tgtgagcttc	cttacataga
11761	agagggggat	gaaggcgagg	aggaagaggg	cgagtacttg	gaagactgat	ggcacaaccc
11821	atattttta	ctagatggaa	cagcaagcac	cggatcccgc	aatgcgggcg	gcgctgcaga
11881	accaaccate	coocattaac	tcctcggacg	attggaccca	ggccatgcaa	cgtatcatgg
11941	cattaacaac	tcgcaacccc	gaagccttta	gacagcaacc	ccaggccaac	cgtctatcgg
12001	ccatcatgga	agctgtagtg	ccttcccgct	ctaatcccac	tcatgagaag	gtcctggcca
12061	tcotoaacoc	attaataaa	aacaaagcta	ttcgtccaga	tgaggccgga	ctggtataca
12121	accetetett	agaacgcgtg	gctcgctaca	acagtagcaa	tgtgcaaacc	aatttggacc
12181	gtatgataac	agatgtacgc	gaagccgtgt	ctcagcgcga	aaggttccag	cgcgatgcca
12241	acctgggttc	actaataaca	ttaaatgctt	tcttgagtac	tcagcctgct	aatgtgccgc
12301	gtggtcaaca	ggattatact	aactttttaa	gtgctttgag	actgatggta	tcagaagtac
12361	ctcagagega	agtatatcag	tecaateeta	attacttctt	tcagactagc	agacagggct
12421	tocagacoot	aaatctgagc	caagctttta	aaaaccttaa	aggtttgtgg	ggagtgcatg
12481	ccccaataga	agaaagagca	accgtgtcta	gcttgttaac	tccgaactcc	cgcctattat
12541	tactottoot	agctcctttc	accgacagcg	gtagcatcga	ccgtaattcc	tatttgggtt
12601	acctactaaa	cctgtatcgc	gaagccatag	ggcaaagtca	ggtggacgag	cagacctatc
12661	aagaaattac	ccaagtcagt	cacactttag	gacaggaaga	cactggcagt	ttggaagcca
12721	ctctgaactt	cttgcttacc	aatcootctc	aaaagatccc	tcctcaatat	gctcttactg
12781	cogaggagga	gaggateett	agatatgtgc	agcagagcgt	gggattgttt	ctgatgcaag
12841	aggggggaac	tecgaetgea	gcactggaca	tgacagcgcg	aaatatggag	cccagcatgt
12901	atoccaotaa	ccgacctttc	attaacaaac	tgctggacta	cttgcacaga	gctgccgcta
12961	tgaactctga	ttatttcacc	aatoccatct	taaacccgca	ctggctgccc	ccacctggtt
13021	tctacacaga	cgaatatgac	atoccoacc	ctaatgacgg	atttctgtgg	gacgacgtgg
13021	acadedatat	tttttcacct	ctttctgatc	atcocacoto	gaaaaaggaa	ggcggcgata
13141	gaatgcattc	ttctgcatcg	ctatccaaaa	tcattggtgc	taccgcggct	gagecegagt
13201	ctacaeatca	ttttcctagt	ctaccetttt	ctctacacag	totacotacc	agcgaagtgg
13261	gtagataag	tcgcccgagt	ttaatgggcg	aagaggagta	cctaaacgat	tecttgetca
13321	gragaaraag	agaaaaaaat	ttcccaaaca	atggaataga	aagtttggtg	gataaaatga
13321	gaccggcaag	gacttatgct	caggatcaca	gagacgagcc	toggatcato	gggactacaa
13//1	gtagatggaa	ccgtagacgc	caggaccata	acagacagag	agatettata	toggacgatg
12501	accettcoc	cgatgatagc	agcgtattgg	acttgggtgg	gagaggaagg	ggcaacccgt
13561	ttactcattt	gegeeetege	ttagatagta	tottotaaaa	aaaaataaaa	aagaaaaaac
13631	teaccaaccc	catggcgacg	accotacott	cattettett	tattatctgt	otctaotata
13681	atrarrorar	tcgtgctagg	cadaacaata	gtgtatccgg	agggtcctcc	tccttcgtac
127/1	acyayycyay	tgcagcagca	acsaacasca	acaataataa	aatccccact	ggaggctccc
12001	yayaycycya	cgcgatacct	gcaggcgacg	geggegaege	acagcattcg	ttactcggaa
13061	atagasacta	agtacgatac	caccaggettg	tatctggtgg	acaacaaqtc	ggcggacatt
13001	cattetatas	actatcagaa	traccacare	aacttcttga	ccacggtggt	gcaaaacaat
12001	gettetega	ctacggaagc	carcaccago	accattaact	ttgatgaacg	atcocootoo
13301	gactitacce	taaaaaccat	catgcatact	aacatoccca	acgtgaacga	gtatatgttt
14141	ggcggtcagc	tcaaagcgcg	tatgeatact	tccacaaaaac	ctcctgaggg	tottagagta
14101	agcaacaagc	atgatcataa	ccaacatatt	ctassatacq	agtogttcga	otttacttto
T470T	gacgataatt	acttttcggt	gcaagacact	atcracttra	traacaatro	catcatagac
14221	ccagaaggca	aagtgggcag	accacyact	atattaass	atascattaa	tattaaatta
.T470T	aditacitya	acttcaagtt	acagaacgga	ccacaaacta	agttgatcat	acctaggett
14241	gacactagga	aggccttcca	toctcacate	atattactac	ctaactacaa	agtggacttt
14401	tacacctatg	gtctgagcaa	cattattaca	attacasaca	aacacccatt	ccaagagggt
14401	accgaaagcc	tgtatgagga	tttegge	granatatte	carccctttt	agatatagat
14521	tttaagatet	acagcaagaa	acatcasasa	ggaaacaccc	aagetgetge	agaagctaaa
14201	gettatyaya	ttgccaacga	tccaataaa	gtaataatag	ctactcaaat	caddddadac
14041	gcaaacarag	caacatccgt	taggagtagg	geggeeddeg	tagegadae	atctcassac
147/UL	agttttgeeg	aactcactat	taagggtgtg	gaaccaccac	rcassacar	aanttacaat
14/01	acagagicaa	ataaaatcaa	caageeege	gaaaaagacg	acctttcata	caattatooc
14001	gtgttggaag	aaggagtgcg	ttactaceae	ttactcacca	cctcagatgt	cacctacaga
14881	gaccccgaaa	tctactggtc	cccccyyaca	atratrosor	atcetatese	tttccactcc
14941	geggageagg	tetaetggte	gettecagae	acyacycayy	ttatagagat	cttttcaaac
12001	actagacaag	tcagtaacta	tatatasta	ggtgcagage	acceptacea	ctcactteca
15061	agcttctaca	acgaacaagc	LyLytactcc	cagcagcucc	gecayeecae	gaggagaatt
15121	cacgtcttca	accgctttcc	Lgagaaccag	accidated	greegeegge	attacass
15181	accaccgtca	gtgaaaacgt	teetgetete	acagatcacg	ggacccctgcc	gregeage
15241	agtatccggg	gagtccaacg	tgtgaccgtt	actgacgcca	gacgccgcac	tttata
15301	gtgtacaagg	cactgggcat	agregeaceg	cgcgtccttt	caageegeae	cttctddddd
15361	aaaaaaaaaa	atgtccgttc	ttatctcgcc	cagtaataac	accountage	grergegege
15421	tcccagcaag	atgtacggag	gcgcacgcaa	acgutetace	caacateeeg	cacacacacac
15481	cgggcatttt	cgcgctccat	ggggtgccct	caagggccgc	actogogtto	gaaccaccgt
15541	cgatgatgta	atcgatcagg	cggttgccga	cgcccgtaat	catactccta	cigogectae
						•

FIG. 28A-4

15601	atctactoto	gacgcagtta '	ttgacagtgt	agtggctgac	gctcgcaact	atgctcgacg
15001	teerare	cgaaggcgca	ttaaaaaaa	traccraret	accactorca	tacaaacaac
TOOOT	taagagccgg	cyaagycyca	ctyccagacy	ccaccgagec	b	
15721	aagagctctg	ctacgaagag	ctagacgcgt	ggggcgaaga	gecatgetta	gggcggccag
15781	acotocaoct	tcgggcgcca	gcgccggcag	gtcccgcagg	caagcagccg	ctgtcgcagc
158/1	anchactatt	gccgacatgg	cccaatcgcg	aagaggcaat	gtatactggg	tgcgtgacgc
17041	beesses	caacgtgtac	coatacages	coatcocct	cacacttaga	agatactgag
TOAUT	tgccaccggt	Caacytytat	ccgcgcac			agacacogas
15961	cagtctccga	tgttgtgtcc	cageggegag	gatgtccaag	cgcaaataca	aggaagaaac
16021	gctgcaggtt	atcgcacctg	aagtctacgg	ccaaccgttg	aaggatgaaa	aaaaaccccg
16081	caaaatcaaq	cgggtaaaaa	aggacaaaaa	agaagaggaa	gatogcoatg	atgggctggc
10001		cgcgagtttg	CCCCSCCCCC	acacatacaa	tancatanac	gcaaagttcg
10141	gyagurugug	cycyaycccy		-Literson		geaccactec
16201	acatgtgttg	agacctggaa	ctteggtggt	Ciciacacce	ggcgagcgcc	caagegeeae
16261	ttttaagcgt	tcctatgatg	aggtgtacgg	ggatgatgat	attcttgagc	aggcagctga
16321	ccgattaggc	gagtttgctt	atggcaagcg	tagtagaata	aatcccaagg	atgaaacagt
16381	atcestacce	ttggatcatg	gaaatcccac	ccctagtctt	aaaccootca	ctttqcagca
10301		gtaactccgc	gaacoccat	tasacacasa	aataaaatt	totateceae
10441	agigulacic	graactccgc	gaacaggege		ggtgaagatt	agetassart
16501	tatgcaactg	atggtgccca	aacgccagaa	gttggaggac	gttttggaga	aaytaaaayt
16561	ggatccagat	attcaacctg	aggttaaagt	gagacccatt	aagcaggtag	cgcctggtct
16621	gggagtacaa	actgtagaca	ttaaaattcc	cactgaaagt	atggaagtgc	aaactgaacc
16681	cacasaacct	actgccacct	ccactgaagt	gcaaacggac	ccatogatgc	ccatgcctat
10001	tecadageee	gccgtcggtc	coactgaage	atocoraços	aantacooto	carcaantct
10/41	Lacaactgac	geegeeggee	ccactcgaag	L-LL-LL-L	aagtatggtt	2444444
16801	gttgatgccc	aactatgtcg	tacacccatc	tattatteet	acteetggtt	accyayycac
16861	tcgctactat	cgcagccgaa	acagtacttc	ccgccgtcgc	cgcaagacac	ctgcaaatcg
16921	cagtcgtcgc	cgtagacgca	caaqcaaacc	gattcccggc	gccctggtgc	ggcaagtgta
16001	ccacaataat	agtgcggaac	ctttgacact	accaeataca	cottaccatc	ctagtatcat
10901	ccgcaacggc	atgttgccgc	tecatactta	googlegeg	cctcacttat	caccttcaca
1/041	cacttaatca	argriguege	Lycettetty	cagacacage	ccccacccgc	ttaaaaaaaa
17101	ttcccatcac	tggttaccga	ggaagaaact	cgcgccgtag	aagagggarg	rrggggcgcg
17161	gaatgcgacg	ctacaggcga	cggcgtgcta	tccgcaagca	attgcggggt	ggttttttgc
17221	cagccttaat	tccaattatc	gctgctgcga	ttggcgcaat	accaggcata	gcttccgtgg
17281	coattcance	ctcgcaacga	cattgacatt	ggaaaaaaa	aaaacotata	aataaaaaat
17201	cygttcagge	ctgacactcc	taataatata	agtatatttt	cttagagatg	gaagacatca
1/341	acaatggact	Cigacacicc	Lygiacigig	actatgttt	cccagagacg	teeseeses
17401	atttttcatc	cttggctccg	cgacacggca	cgaagccgta	catgggcacc	Lygageyaca
17461	teggeacgag	ccaactgaac	gggggcgcct	tcaattggag	cagtatctgg	agcgggctta
17521	aaaattttgg	ctcaaccata	aaaacataco	ggaacaaagc	ttggaacagc	agtacaggac
17591	agggggttag	aaataaactt	aaagaccaga	acttccaaca	aaaagtagtc	gatgggatag
17701	aggegeetag	caatggagtg	atagaccaga	ataaccacc	tatacease	aanataaana
1/641	cttccggtat	caatggagtg	grayarrryg	ctaactagge	cgcgcagaaa	aagattaata
17701	gtcgtttgga	cccgccgcca	gcaaccccag	gtgaaatgca	agtggaggaa	gaaatteete
17761	cgccagaaaa	acgaggcgac	aagcgtccgc	gtcccgattt	ggaagagacg	ctggtgacgc
17821	gcgtagatga	accgccttct	tatgaggaag	caacgaagct	tggaatgccc	accactagac
17881	coatacccc	tatggccacc	ggggtgatga	aaccttctca	attacatcaa	cccgtcacct
17001	teactiteaa	ccctcctcct	actactecta	ctatacccac	ttctaacct	atcactaccc
1/941	Lygalliged	CCCCCCCC	getgetactg		tactactes	antagagaga
18001	cgaaaccagt	cgccgtagcc	aggtcacgtc	eegggggege	teetegteea	aatycacact
18061	ggcaaaatac	tctgaacagc	atcgtgggtc	taggcgtgca	aagtgtaaaa	cgccgtcgct
18121	gcttttaatt	aaatatggag	tagcgcttaa	cttgcctatc	tgtgtatatg	tgtcattaca
18181	caccatcaca	gcatcagagg	aaaaaaaaaaaa	gaggtcgtgc	atcaacacta	agttactttc
10201	2000000000	ccccatcgat	actaccces	taggestaca	tocacateoc	cogacaggat
10241	aayatyytta	t	getgetetaa		acacacacac	cacctacttc
18301	gcttcggagt	acctgagtcc	gggtctggtg	cagttegeec	yegecacaga	Cacciacic
18361	aatctgggaa	ataagtttag	aaatcctacc	gtagcgccga	cccacgatgt	gaccaccgat
18421	cataaccaac	ggctcatgtt	gcgcttcgtg	cccgttgacc	gggaggacaa	tacatactct
18481	tacaaagtgc	ggtacaccct	aaccataaac	gacaacagag	tgctggatat	ggccagcacg
105/1	ttatttaaa	ttaggggcgt	nttonacana	gatacaatt	ttaaacccta	ttctggtacg
10041	Licitiyaca	LLaggggcgL	griggadaga	ggccccagcc	atasataatt	ccctggcacg
18601	gcttacaact	ccctggctcc	taaaggeget	ccaaatgcat	Ctcaytyytt	gyacaaggya
18661	gttacaagca	ctggcctagt	ggacgacggc	aatactgatg	atggggaaga	agccaaaaaa
18721	gcaacataca	cttttggtaa	tgctccagta	aaagccgagg	ctgaaatcac	aaaagacgga
18781	ttaccaataa	gcttggaagt	ttcaactgaa	ggtcctaaac	caatctatoc	tgataagctt
10041	tateaceae	aacctcaagt	CCCCCC Sar	acttmmartm	acctadacdd	aaaaacccaa
T004T	Laccagecag	aaccccaagt	gggagacgaa	actiggateg	22000	accetattt
18901	gagtatggag	ggagggttct	taaacctgaa	actaaaatga	aaccctgcta	cygatettt
18961	gctaaaccta	ctaatattaa	aggaggtcag	gcaaaggtaa	. aaccaaaaga	agacgatggc
19021	actaacaaca	tcgagtatga	cattgacatg	aacttctttg	acttaagatc	acaaagatca
19091	caactcaaac	ctaaaattgt	aatgtatgca	gaaaatgtgg	acctggaatg	tccagatact
T200T	gaactcaaac	acaaacctgg	antetanast	20200000000000000000000000000000000000	anarcaatot	tanacasasa
19141	catginging	acaaaccugg	agicicagat	golagilolg	agaccaattt	- Lygacaacay
19201	tctatgccca	acagacccaa	ctacattggc	ttcagagata	acticategg	acttatgtac
19261	tataacagta	ctggcaacat	gggggtactg	gctggccaag	cgtctcagtt	gaatgcagtg
19321	attaacttac	aggacagaaa	cacagaactg	tcttaccaac	: tcttgcttga	ctctctgggc
19321	2-3-2-2-3-	gatactttag	catotogaat	caggetataa	acagttatoa	tcctgatgta
10444	gatayaatta	aaaatcatgg	tataasaat	reacttrors	actattott	tecattanet
エフセゼー	cycyclatey	addaccatyy	-y cygaayat	gualticood		gggat

40504						
TAPOT	ggtgtcggtc	cgcgaacaga	tagttacaag	gagattaagc	caaatggaga	ccaatctact
19561	togacaaato	tagacccaac	taggaggagt	gaacttgcta	agggaaatcc	atttgccatg
10001	cggacaaacg	cagacccaac	cggcagcagc	gaactegeta	agggaaatoo	
19621	gaaattaacc	ttcaagccaa	tctatggcga	agtttccttt	attccaatgt	ggctctatat
19681	ctcccagact	cotacaasta	caccccaticc	aatotcacto	ttccagaaaa	caaaaacacc
10001	Cccccagacc	Caracaara	caccocgacca	aacgccaccc	Leccagaaaa	
19741	tacgactaca	tgaacgggcg	ggtggtgccg	ccatctctag	tagacaccta	tgtgaacatt
19801	antacceant	aatetetaaa	taccatagae	aatotcaacc	cattcaacca	ccaccotaac
13001	garaccagar	ggcccccgga	caccacaaac	aacgccaacc	Caccoaacca	to the same trans
19861	gctggcttgc	gttaccgatc	catgcttctg	ggtaacggac	gttatgtgcc	tttccacata
10021	gaagtgggtg	aaaaattott	cactattess	pacetactac	ttctcccagg	ctcctacact
19981	tatgagtgga	actttaggaa	ggatgtaaac	atggttctac	agagttccct	cggtaacgac
					acctctatgc	
20101	cccatggctc	acaacaccgc	ttccaccctt	gaagccatgc	tgcggaatga	caccaatgat
					accccattcc	
20221	accaatattc	ccatttccat	teettetege	aactgggcgg	ctttcagagg	ctggtcattt
					gatttgaccc	
20341	tattctggtt	ctattcccta	cctagatagt	accttctacc	tgaaccacac	ttttaagaag
20401		tetttenata	ttaaataaaa	tagaatagaa	atgacaggtt	actatotoot
20461	aacgaatttg	aaataaagcg	cactotogat	ggcgaaggct	acaacgtagc	ccaatgcaac
20221	atgaccaaag	actggttett	ggcacagacg	CLEGECAACL	acaacatcgg	Clattayyyt
20581	ttctacattc	cagaaggata	caaagatcgc	atotattcat	ttttcagaaa	cttccagccc
20641			tesestesst	+	tassagaat	ccccataccc
7004T	atgageagge	aggiggilga	cgaggicaai	Lacaaagaci	tcaaggccgt	cyccataccc
20701	taccaacaca	acaactctgg	ctttataaat	tacatooctc	cgaccatgcg	tcaaqqtcaa
20761						
20761	ccctatcccg	ctaactatcc	ctatccactc	attggaacaa	ctgccgtaaa	tagtgttacg
20821	cagaaaaagt	tettatataa	cagaaccatg	tggcgcatac	cgttctcaag	caacttcato
20881	tctatgggag	cccttacaga	cttgggacag	aacatgctct	atgccaactc	agctcatgct
					ccctgcttta	
21001	gaagttttcg	acgtggtcag	agtgcatcag	ccacaccgcg	gcatcatcga	ggcagtctac
21061	atagatagaa	cattatagas	contagonet	accacutaan	aagcttcttg	cttcttccaa
21001	Cigigiacac	cyccccggc	cggcaacgcc	accacgcaag	aagccccccg	CEECELGCAA
21121	acagcagctg	caaccatggc	ctgcggatcc	caaaacggct	ccagcgagca	agagctcaga
21191	gggattatas	aagacotggg	ttacaaacca	tatttttaa	gaacctttga	taaccacttc
21101	gccattgccc	aagacccggg	ctgcggacca	cacccccgg	gaacettega	caagegeeee
21241	ccggggttca	tggcccccga	taagctcgcc	tgtgccattg	taaatacggc	cggacgtgag
21301	2000000000	aggactggtt	agetttaggt	tagaaccac	gttctaacac	ctactacett
21361	tttgatcctt	ttagattctc	ggatgatcgt	ctcaaacaga	tttaccagtt	tgaatatgag
21421	ggteteetge	geegeagege	LCLLyctacc	aaggaccggc	gtattacgct	ggaaaaattt
21481	acccagaccg	tacagaaccc	ccattctacc	acctacaac	ttttctgctg	catottcctt
21541		teeneteese	tasaaataaa	atamagana	200002002	gaaattggta
					accccaccat	
21601	actogagtoc	caaacaacat	gcttcattct	cctaaagtcc	agcccaccct	gtgtgacaat
					attttcgctc	
21721	cacatcgaaa	gggccactgc	gttcgaccgt	atggatgtgc	aataatgatt	catotaaaca
01701		555000050				
7T/8T	acgtgttcaa	taaacagcac	tttattttt	acatgtateg	aggctctgga	CLACTTACTE
21841	atttacaagt	cgaatgggtt	ctgacgagaa	tcagaatgac	ccgcaggcag	tgatacgttg
					ccaacttggg	
21961	t.cgggcagga	totcactcca	cagetttetg	atcaactaca	aagctcccag	caggtcagga
22021						
		tyaaattata	attraquacca			accetococc
22081				gracectag	cgcgagagtt	gcggtacacc
	ddattdcadc					gcggtacacc
		actgaaacac	catcagcgac	ggatgtctta	cgcttgccag	gcggtacacc cacggtggga
	tctgcaatca	actgaaacac tgcccacatc	catcagcgac cagatcttca	ggatgtctta gcattggcaa	cgcttgccag tgctgaacgg	gcggtacacc cacggtggga ggtcatcttg
22201	tctgcaatca	actgaaacac tgcccacatc	catcagcgac cagatcttca	ggatgtctta gcattggcaa	cgcttgccag tgctgaacgg	gcggtacacc cacggtggga ggtcatcttg
	tctgcaatca caggtctgcc	actgaaacac tgcccacatc tacccatggc	catcagcgac cagatcttca gggcacccaa	ggatgtctta gcattggcaa ttaggcttgt	cgcttgccag tgctgaacgg ggttacaatc	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg
22261	tctgcaatca caggtctgcc gggatcagta	actgaaacac tgcccacatc tacccatggc tcatcttggc	catcagcgac cagatcttca gggcacccaa ctgatcctgt	ggatgtctta gcattggcaa ttaggcttgt ctgattcctg	cgcttgccag tgctgaacgg ggttacaatc gatacacggc	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa
22261	tctgcaatca caggtctgcc gggatcagta	actgaaacac tgcccacatc tacccatggc tcatcttggc	catcagcgac cagatcttca gggcacccaa ctgatcctgt	ggatgtctta gcattggcaa ttaggcttgt ctgattcctg	cgcttgccag tgctgaacgg ggttacaatc gatacacggc	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa
22261 22321	tetgcaatca caggtetgce gggatcagta gcatcatatt	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct	ggatgtctta gcattggcaa ttaggcttgt ctgattcctg ttactaccct	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag
22261 22321 22381	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaactggtt	catcagegae cagatettea gggcaeceaa ctgateetgt ctgetggget agetgegeag	ggatgtetta gcattggcaa ttaggettgt ctgatteetg ttactacect ccggcatcat	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcacacagca	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca
22261 22321 22381	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaactggtt	catcagegae cagatettea gggcaeceaa ctgateetgt ctgetggget agetgegeag	ggatgtetta gcattggcaa ttaggettgt ctgatteetg ttactacect ccggcatcat	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcacacagca	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca
22261 22321 22381 22441	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggcta	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac	catcagegac cagatettea gggcacceaa ctgateetgt ctgetggget agetgegeag acttetgece	ggatgtetta gcattggcaa ttaggettgt etgatteetg ttactaceet eeggeateat eageggtttt	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcacacagca gggtgatttt	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca ggttcgctcg
22261 22321 22381 22441 22501	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggcta ggattctcct	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac tcaaggctcg	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct agctgcgcag acttctgccc ttgtccgttc	ggatgtetta gcattggcaa ttaggettgt ctgattectg ttactaccet ceggcateat cageggtttt tegetggcca	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcacacagca gggtgatttt catccatctc	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca ggttcgctcg gataatctgc
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22261 22321 22381 22441 22501 22621 22681 22741 22801 22921 22981 23041 23161 23161 23221	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggta ggattctcct tccttctgaa ccatgaggcc gaatgtatca aaagttaact tgttcgtgct ttctccatca ctaatcggat acttctcaa ccactgcagta atcttctccaa ccactgcagta tctctccatca ctaatcggat atcttctccat atggggacat tcggtagaag	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac tcataggctcg tcataatatt acaacgcaca ttccctgcag ggatgcctcg gctcaggcat gcagacacat tcttaacagt tgcttctttt caagttggc gtttggtctt ccggagacac aacctgaccc aacctgaccc	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct agctgcgcag acttctgccc ttgtccgttc gccatgcaag gcctgtacat aaatcttccc gtgctcctcg tagtttaaaa cacttccatg gcagcagca gccatccttc ctcttctct cgtggttc	ggatgtetta gcattggcaa ttaggettgt ctgattectg ttactaccet ccggcatcat cagcggtttt tcgctggcca cacttcaget tcccaattat atcategtge ttcacgtact gaggttetaa cctttetece gctacctatag tcaacgaget tctattetegg gacgttteteg gaggtttete	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcaccatctc tgccctcata ggtggcgat tcagtgtctt ggtgacagat gttcgttatc aagcagacac ccagagggcg tgtcttgact gtatcgagg tgtcttgact gtatcggggg tcaccattac	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgag gcgggcgtca ggttcgctcg gataatctgc atcattgcag ctgagaaaaa gtgactagtg gcgcttgtat cagcctgtac caggggcaag atctttggcg gtagctgaaa gatgtctaga gagatgactg gcaactgactg cagaggtgga
22261 22321 22381 22441 22501 22621 22681 22741 22801 22921 22981 23041 23161 23161 23221	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggta ggattctcct tccttctgaa ccatgaggcc gaatgtatca aaagttaact tgttcgtgct ttctccatca ctaatcggat acttctcaa ccactgcagta atcttctccaa ccactgcagta tctctccatca ctaatcggat atcttctccat atggggacat tcggtagaag	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac tcataggctcg tcataatatt acaacgcaca ttccctgcag ggatgcctcg gctcaggcat gcagacacat tcttaacagt tgcttctttt caagttggc gtttggtctt ccggagacac aacctgaccc aacctgaccc	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct agctgcgcag acttctgccc ttgtccgttc gccatgcaag gcctgtacat aaatcttccc gtgctcctcg tagtttaaaa cacttccatg gcagcagca gccatccttc ctcttctct cgtggttc	ggatgtetta gcattggcaa ttaggettgt ctgattectg ttactaccet ccggcatcat cagcggtttt tcgctggcca cacttcaget tcccaattat atcategtge ttcacgtact gaggttetaa cctttetece gctacctatag tcaacgaget tctattetegg gacgttteteg gaggtttete	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcaccatctc tgccctcata ggtggcgat tcagtgtctt ggtgacagat gttcgttatc aagcagacac ccagagggcg tgtcttgact gtatcgagg tgtcttgact gtatcggggg tcaccattac	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgag gcgggcgtca ggttcgctcg gataatctgc atcattgcag ctgagaaaaa gtgactagtg gcgcttgtat cagcctgtac caggggcaag atctttggcg gtagctgaaa gatgtctaga gagatgactg gcaactgactg cagaggtgga
22261 22321 22381 22441 22501 22621 22621 22861 22921 22981 23041 23161 23221 23281	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggcta ggattctcct tccttctgaa ccatgaggcc gaatgtatca acatgaggcc ttctccatca ctaatcggat atctctcaa ccactgcta atctctcaa ccactgcta atcgggacat ttcggggacat tcggtagaag ggcgattgcg	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac tcataggctcg tcataatatt acaacgcaca ttccctgcag ggatgcctcg gctcaggcat gcagacacat tcttaacagt tgcttcttt caagttggctct gcttgggcaca tcgtgggcaca tcgtgggcaca tcaggcaca	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct agctgcgcag acttctgccc ttgtccgttc gccatgcaag gcctgtacat aaatcttccc gtgctcctcg tagtttaaaa cacttccatg gcaggcagca gccatccttc ctcttctct cgtgggttg cacacggcga gtccacctg	ggatgtetta gcattggcaa ttaggettgt ctgattectg ttactaccet ccggcatcat cagcggtttt tcgctggcca cacttcaget tcccaattat atcategtge ttcacgtact gaggttetaa cctttetece getectttag tcaacgaget tctatteggg tcaacgaget tctattetegg tcaacgaget tcttetteggg tagtgtttc gaaggeggat	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcacatctt catccatctt tgccctcata ggtggcgat tcagtgtctt ggtgacagat gttcgttatc aagcagacac ccagagggtc gcacgggcgg tgtcttgact gtatcggag tcaccattac tcttcggggg gactggcaga	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca ggttcgctcg gataatctgc atcattgcag ctgagaaaaa gtgactagtg gcgcttgtat cagcctgtac caggggcaag atctttggcg gtagctgaaa gatgatgtgcag gagatgactg cagaggtgga caactgactg cagaggtgga accccttccg
22261 22321 22381 22441 22501 22621 22621 22861 22921 22981 23041 23161 23221 23281	tctgcaatca caggtctgcc gggatcagta gcatcatatt gacctgctcg ttgttggcta ggattctcct tccttctgaa ccatgaggcc gaatgtatca acatgaggcc ttctccatca ctaatcggat atctctcaa ccactgcta atctctcaa ccactgcta atcgggacat ttcggggacat tcggtagaag ggcgattgcg	actgaaacac tgcccacatc tacccatggc tcatcttggc gcttgaaagc aaaactggtt tttgcaccac tcataggctcg tcataatatt acaacgcaca ttccctgcag ggatgcctcg gctcaggcat gcagacacat tcttaacagt tgcttcttt caagttggctct gcttgggcaca tcgtgggcaca tcgtgggcaca tcaggcaca	catcagcgac cagatcttca gggcacccaa ctgatcctgt ctgctgggct agctgcgcag acttctgccc ttgtccgttc gccatgcaag gcctgtacat aaatcttccc gtgctcctcg tagtttaaaa cacttccatg gcaggcagca gccatccttc ctcttctct cgtgggttg cacacggcga gtccacctg	ggatgtetta gcattggcaa ttaggettgt ctgattectg ttactaccet ccggcatcat cagcggtttt tcgctggcca cacttcaget tcccaattat atcategtge ttcacgtact gaggttetaa cctttetece getectttag tcaacgaget tctatteggg tcaacgaget tctattetegg tcaacgaget tcttetteggg tagtgtttc gaaggeggat	cgcttgccag tgctgaacgg ggttacaatc gatacacggc cggtataaaa tcaccatctc tgccctcata ggtggcgat tcagtgtctt ggtgacagat gttcgttatc aagcagacac ccagagggcg tgtcttgact gtatcgagg tgtcttgact gtatcggggg tcaccattac	gcggtacacc cacggtggga ggtcatcttg gcagtgcagg tctcatgaaa catcccgcag gcgggcgtca ggttcgctcg gataatctgc atcattgcag ctgagaaaaa gtgactagtg gcgcttgtat cagcctgtac caggggcaag atctttggcg gtagctgaaa gatgatgtgcag gagatgactg cagaggtgga caactgactg cagaggtgga accccttccg

FIG. 28A-6

			•			<b>*</b>
23401	gtgttctcct	aggcagagaa	acaacagaca	tggaaactca	gccattgctg	ccaacacege
23461	cacgagtgcc	atcacatctc	qtcctcaqcq	acqaqqaaaa	ggagcagagc	ttaagcattc
22521	0200000000	testaceace	acctetacee	tanaanataa	ggaggtcgac	gcatctcatg
23521	caccycccag	tectgecace	acciccaccc	tagaagataa	9949940944	eaccetate
23581	acatgcagaa	taaaaaagcg	aaagagtctg	agccagacat	cgaacaagac	cegggerary
23641	tgacaccggt	ggaacacgag	gaagagttga	aacgctttct	agagagagag	gatgaaaact
23701	CCCC2222C2	gcaagcggat	aactatcacc	aagatgctgg	aaatagggat	cagaacaccg
23/01	ycccaaaaca	gcaugeggae		auguegeegg		agaaataaa
23761	actacctcat	agggcttgac	ggggaagacg	cgctccttaa	acatctagca	agacagucac
23821	tcatagtcaa	ggatgcatta	ttggacagaa	ctgaagtgcc	catcagtgtc	gaagagctca
23881	accacacata	cgagettaac	ctattttcac	ctcatactcc	ccccaaacgt	cagccaaacg
23001	geegegeeta	cgageetaae		tttataaaa	ttttaatata	casasatsc
23941	gcacctgcga	gccaaatect	egettaaact	LLLatecaye	ttttgctgtg	ccagaagtac
24001	tggctaccta	tcacatcttt	tttaaaaatc	aaaaaattcc	agtctcctgc	cgcgctaatc
24061	deseccacae	coatocccta	ctcaatctgg	gacctggttc	acgcttacct	gatatagett
24121	generages	agttagaaaa	atottorage	atctagacee	taatgagact	caaaccacaa
24121	ccccggaaga	ggttccaaag	accurage	gicigggcaa	caacyayacc	cygyccycaa
24181	atgctctgca	aaagggagaa	aatggcatgg	atgagcatca	cagcgttctg	gtggaattgg
24241	aaggcgataa	tgccagactc	gcagtactca	agcgaagcgt	cgaggtcaca	cactttgcat
24301	acccccctat	caacctgccc	cctaaagtca	tgacggccgt	catggaccag	ttactcatta
24361	20000000	tacactttca	gaagaataa	atracceara	tgcctgtgat	gagggtaaac
24301	agegegeaag	LUCCULLUCA	gaagacacgc	atyacccaga		
24421	cagtggtcag	tgatgagcag	ctaacccgat	ggctgggcac	cgactctccc	cgggatttgg
24481	aagagcgtcg	caagettatg	atggccgtgg	tgctggttac	cgtagaacta	gagtgtcttc
2/5/1	aggatttett	taccoattca	gaaaccttgc	gcaaactcga	agagaatctg	cactacactt
24341	bbassassass			acatatataa	agegeeets	accaacctcc
24601	ttagacacgg	ctttgtgcgg	caggcargca	ayatatttaa	cgtggaactc	accaaccigg
24661	tttcctacat	gggtattctg	catgagaatc	gcctaggaca	aagcgtgctg	cacagcaccc
24721	ttaaggggga	agecegeegt	gattacatcc	gcgattgtgt	ttatctctac	ctgtgccaca
24701	catagggga	caacataaat	atataacaac	aatotttaga	agaacagaac	ctgaaagagc
24/01	cycyycadac	cygcacyggc	t-t-t-t-	ttatataaaa	agaacagaac	2222222
24841	taaacaagct	cttacagaaa	tetettaagg	Licigiagae	agggttcgac	gagegeaceg
24901	tcgcttccga	cctggcagac	ctcatcttcc	cagagcgtct	cagggttact	ttgcgaaacg
24961	gactgcctga	ctttatgage	cagagcatgc	ttaacaattt	tcgctctttc	atcctggaac
25021	gatagaatat	cetacecaca	acctactaca	cactoccctc	cgactttgtg	cctctcacct
25021	getteggtat	cctgcccgcc	accigcigcg	ttt	-tttate	eeeeeeeeee
25081	accgcgaatg	cccccgccg	ctatggagte	actgctacct	gttccgtctg	gecaactace
25141	tctcctacca	ctcggatgtg	atcgaggatg	tgagcggaga	cggcttgctg	gagtgtcact
25201	accactacaa	tetatacaca	ccccaccoot	ccctagettg	caacccccag	ttgatgagcg
25261	googoogoat	aatagggagg	tttgaattgg	aaggggggg	cagccaaggc	datdddtctt
23201	adacccagac	aatayytatt	tttgaattgt	aaggeeeeag	cagecaagge	******
25321	ctcctgggca	aagtttaaaa	ctgaccccgg	gactgtggac	ctccgcctac	Ligitgeaagi
25381	ttgccccgga	agattaccac	ccctatgaaa	tcaagttcta	tgaggaccaa	tcacagcctc
25441	cuaaaaccaa	actttcggcc	tocotcatca	cccagagaac	aattctggcc	caattgcaag
25501	castagoogu	atcocccaa	caatttctac	tassssaaa	taagggggtc	taccttgacc
2550I	CCatCCaaaa	accecyceaa	gaacttttat	Lyaaaaaggg	taagggggcc	
25561	cccagaccgg	cgaggaactc	aacacaaggt	tccctcagga	tgtcccaacg	acgagaaagc
25621	aagaagttga	aggtgcagcc	gccgccccca	gaagatatgg	aggaagattg	ggacagtcag
25681	acadaddaad	содаддадда	ggacagtetg	gaggacagtc	tggaggaaga	cagtttggag
25001	gougugguug	0994994994	33-0030003	gaggaragee	CCGaCaaaca	attatected
25/4I	gaggaaaacg	aggaggcaga	ggaggragaa	gaagtaaccg	ccgacaaaca	gecaececeg
25801	gctgcggaga	caagcaacag	cgctaccatc	tccgctccga	gtcgaggaac	ccggcggcgt
25861	cccagcagta	gatgggacga	gaccggacgc	ttcccgaacc	caaccagcgc	ttccaagacc
25921	autaagaagg	atcoocaooo	atacaagtcc	taacaaaaac	ataagaatgc	catcatctcc
25521	5900090099	2009900999	caacatatco	ttaaaaaaaa	gctacttgct	attocarcat
7239T	tgettgeatg	agracagaga	caacacaccc	Licacycygc	gccaccegcc	accccaccac
26041	ggggtgaact	ttccgcgcaa	tgttttgcat	tactaccgtc	acctccacag	CCCCtactat
26101	agccagcaaa	tcccggcagt	ctcgacagat	aaagacagcg	gcggcgacct	ccaacagaaa
26161	accaucaucu	gcagttagaa	aatacacaac	aagtgcagca	acaggaggat	taaagattac
26221	accugacy	accade contract	coccasastt	aagagatgaa	atctttccaa	ccctatatac
20221	agecaacgag	Ccagcycaaa	cccyayaycc	aayaaaccyy	attettetta	t-t-t-
26281	catcttccag	cagagtcggg	gccaagagca	ggaactgaaa	ataaaaaacc	gatetetgeg
26341	ttcgctcacc	agaagttgtt	tgtatcacaa	gagcgaagat	caacttcagc	gcactctcga
26401	adacaccasa	getetettea	acaagtactg	cacactaact	cttaaagagt	aggcagcgac
20101	ggacgccgag		aaaaattaaa	testestes	catgagtaaa	gaattege
20401	egegettatt	Caaaaaaggc	gygaattata	teatectega	cacgagcaaa	bassassass
26521	cgccttacat	gtggagttat	cagccccaaa	tgggattggc	ggcaggcgcc	teeeaggaet
26581	actccacccd	catgaattgg	ctcagcgccq	ggccttctat	gatttctcga	gttaatgata
26641	tacgcgccta	сспавассва	atacttttog	aacagtcagc	tcttaccacc	acgccccccc
26241	22222222	+ccc=~	taaccaaaaa	ccctactctc	ccaggaaagt	cccactages
20/UI	aacaccttaa	Luccagaaal	- cggcccgccg	ccccagigca		gange to the
26761	ccactgtatt	acttcctcga	gacgcccagg	ccgaagtcca	aatgactaat	acadaracac
26821	agttagcggg	cooctccacc	ctatgtcgtc	acaggeeteg	gcataatata	aaacgcctga
26881	tratrarara	ccmammatate	cadctcaacc	acqagtcggt	gageteteeg	cttootctac
7000T			attoccocc	~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~~	ttccttcacc	Cotogtopas
20941	gaccagacgg	aatctttcag	arraceaace	gcgggagacc	ttccttcacc	
27001	ctgttctgac	tttggaaagt	tegtettege	aaccccgctc	gggcggaatc	gggaccgttc
27061	aatttotooa	ggagtttact	ccctctatct	acttcaaccc	cttctccgga	tctcctgggc
27121	actaccocca	coanttrata	CCGaacttca	acconstrac	cgagtcagtg	gacggctacg
07101	actactigga		ctancetcty	tagagacag	catctacacc	actor-co-
2/181	attgatgtct	ggrgacgcgg	cryaycratc	reggetgega	catctagacc	actycegeeg
27241	ctttcgctgc	tttgcccggg	aactcattga	gttcatctac	ttegaactee	ccaaggatca

27301	ccctcaaggt	ccaacccaca	gagtgcggat	tactatcgaa	ggcaaaatac	actctcgcct
27301	gcaacgaatt	theteres	gagogoggat	ratcracco	gaccaddaaa	acaccaccot
2/361	gcaacgaatt	tteteceage	ggcccgcgcc	gategagega	gaccagggaa	-tattatata
27421	ttccatctac	tgcatttgta	atcaccccgg	attgcatgaa	agcetttget	geceracycy
27481	tactgagttt	aataaaaact	gaattaagac	tctcctacgg	actgccgctt	cttcaacccg
275/1	gattttacaa	ccadaadaac	gaaacttttc	ctotcotcca	ggactctgtt	aacttcacct
7/24T	ttcctactca	ccagaagaac	~~*	tagagagatt	ttccacaacc	attttcccta
27601	ttcctactca	caaactagaa	gercaaegae	Lacacegett	Luccayaagu	accccccca
27661	ctaatactac	tttcaaaacc	ggaggtgagc	tccaaggtct	tcctacagaa	aacccttggg
27721	tggaagcggg	ccttotagto	ctaggaattc	ttacaaataa	gcttgtgatt	attctttgct
27701	acctatacac	accttacttc	actttcctac	tagtattata	gtattggttt	aaaaaatggg
2//01	acctatacac	accitigette			godooggood	aaaattaaa
27841	gcccatacta	gtcttgcttg	Ettlactic	gerrraggaa	cogggeeer	ccaaccacga
27901	tccatgtcta	gacttcgacc	cagaaaactg	cacacttact	tttgcacccg	acacaagccg
27961	catctgtgga	attettatta	agtgcggatg	ggaatgcagg	tccgttgaaa	ttacacacaa
20021	taacaaaacc	tagaacaata	ccttatccac	cacatgggag	ccaggagttc	cccactccta
20021	cactgtctct	-t	atasaaatta	categgggt	actaacaaca	ctttcatttt
2808T	cactgtctct	gcccgaggcc	Cigacyguic	caccegeace	agtaataata	
28141	ttctgaaatg	tgcgatctgg	ccatgttcat	gagcaaacag	tattetetat	ggeeteetag
28201	caaggacaac	atcotaacgt	tctccattgc	ttattgcttg	tgcgcttgcc	ttcttactgc
20261	tttactgtgc	gtatgcatac	acctocttot	aaccactcoc	atcaaaaaco	ccaataacaa
20201	agaaaaaaatg	anthonosta	tttatattt	cacacatoo	ttctcttaca	teteteatat
28321	agaaaaaatg	Colladout	LLCLIGLLCA	cagacatgge	-t-totaca	-to-coopeac
28381	ttgtcagcat	tgtcactgcc	gctcacggac	aaacagtcgt	ctctatccct	Ctaggacata
28441	attacactct	cataggaccc	ccaatcactt	cagaggtcat	ctggaccaaa	ctgggaagcg
28501	ttgattactt	tratataatc	tocaacaaaa	caaaaccaat	aatagtaact	tocaacatac
20501	aaaatcttac	attentions	attaccasac	tttacacccc	ttactattat	aattataaca
7820T	aaaatettae	allyallaal	graycaday	ttacagegg	the	ggccacgaca
28621	gatacagtag	tcaatataga	aattacttgg	ttcgtgttac	ccagttaaaa	accacgaaaa
28681	tgccaaatat	ggcaaagatt	cgatccgatg	acaattctct	agaaactttt	acatctccca
287/11	ccacacccga	спаававаас	atcccagatt	caatgattgc	aattottoca	acaataacaa
20741	tggtgatggc	agtastasta	atatocatoc	ttttatatoc	ttatcactac	aaaaagtttc
2880T	tggtgatgge	accaacaaca	atatycatyc	ttttatatge	atttatt	atagagagat
28861	atcctaaaaa	acaagatctc	ctactaaggc	ttaacattta	atttetttt	acacagecac
28921	ggtttccact	accacattcc	ttatgcttac	tagtcttgca	actctgactt	ctgctcgctc
28981	acacctcact	gtaactatag	gctcaaactg	cacactaaaa	ggacctcaag	gtggtcatgt
20001	cttttggtgg	agaatatatg	acaatooato	otttacaaaa	ccatgtgacc	aacctggtag
29041	cttttggtgg	ayaatataty	acaacggacg	teteeeete	agaggata	adoobggoes
29101	atttttctgc	aacggcagag	acctaaccat	tatcaacgtg	acaycaaacy	acaaayyccc
29161	ctattatgga	accgactata	aaagtagttt	agattataac	attattgtac	tgccatctac
29221	cactccagca	ccccccacaa	ctactttctc	tagcagcagt	gtcgctaaca	atacaatttc
20221	caatccaacc	tttaccacac	ttttaaaaco	cactotoaat	aattctacaa	cttcacatac
29201	caatccaacc	the	tectadaacg	anatananta	accottccaa	tatotattot
29341	aacaatttcc	acttcaacaa	teageattat	egetgeagtg	acaactyyaa	Latitlatici
29401	tgtttttacc	ataacctact	acgcctgctg	ctatagaaaa	gacaaacata	aaggtgatcc
29461	attacttaga	tttgatattt	aatttgttct	tttttttt	atttacagta	tggtgaacac
20521	caatcatggt	acctanaaat	ttcttcttca	ccatactcat	ttgtgcattt	aatotttoco
29321	caaccatggt	acctagaaac	2000000000	coccatatat	aggaggattt	acttactata
2958T	ctactttcac	agcagtagee	acaycaaccc	cayactycat	aggagcaccc	gettettatg
29641	cactttttgc	ttttgttact	tgcatctgcg	tatgtagcat	agtctgcctg	grtattaatt
29701	ttttccaact	tctagactgg	atccttgtgc	gaattgccta	cctgcgccac	catcccgaat
29761	accgcaacca	agatategeg	gcacttetta	gactcatcta	aaaccatgca	ggctatacta
20701	ccaatatttt	taatacogog	gottogottag	actatatass	ccccaactac	ctatactact
29821	ccaatatttt	Egelletatt	getteeteac	getgeeteaa	tb-t	testageact
29881	ccaccagaac	accttagaaa	atgcaaattc	caacaaccgt	ggtcatttct	tgettgetat
29941	cgagaaaaat	cagaaattcc	cccaaattta	ataatgattg	ctggaataat	taatataatc
30001	tgttgcacca	taatttcatt	tttgatatac	cccctatttq	attttggctg	gaatgctccc
30061	aatgcacatg	atcatccaca	adacccadad	gaacacattc	ccctacaaaa	catocaacat
20001	aatytataty	accaccaca	agacccagag	gaacacacco	coctactaca	tactattaat
30121	ccaatagcgc	taatagatta	cgaaagtgaa	ccacaacccc	Cactactece	Lyctattagt
30181	tacttcaacc	taaccggcgg	agatgactga	aacactcacc	acctccaatt	ccgccgagga
30241	tctgctcgat	atggacggcc	acateteaga	acagcgactt	gcccaactac	gcatccgcca
30301	gcagcaggaa	cacacacac	aagageteag	agatotoato	caaattcacc	aatocaaaaa
30301	gcaycayyaa	cycycygcca	aayaycccay	agacyccucc	cadattoaco	atactacaa
3036T	aggcatattc	tgtttggtaa	aacaagccaa	gatatectae	gagarcaccg	Ctactgacta
30421	tegeetetet	tacgaacttg	gcccccaacg	acaaaaattt	acctgcatgg	tgggaatcaa
30481	ccccatagtt	atcacccagc	aaagtggaga	tactaagggt	tgcattcact	gctcctgcga
30541	ttccatcgag	tocacctaca	ccctactasa	gaccctatge	ggcctaagag	acctoctacc
20241		cgcacccaca	ccccgccguu	acttacttca	astracrast	and at at at a
2000T	aatgaattaa	aaaatgatta	alaaadaatC	actiactiga		Laggerery
30661	ttgaaatttt	ctcccagcag	cacctcactt	cccccttccc	aactctggta	LECTABACCC
30721		catactttct	ccatacttta	aaggggatgt	caaattttag	ctcctctcct
	Carregacaa			antanagana		trantmartr
30721	cgttcagcgg	tettestate	tttcttccca		agagtccuuc	
30781	gtacccacaa	tcttcatgtc	tttcttccca	gatyactaay	caacacacac	ttatasacco
30781 30841	gtacccacaa cttcaaccct	tcttcatgtc	atgaagatga	aagcacctcc	caacacccct	ttataaaccc
30781 30841 30901	gtacccacaa cttcaaccct agggtttatt	tcttcatgtc gtctacccct tccccaaatg	atgaagatga gcttcacaca	aagcacctcc aagcccagac	caacacccct ggagttctta	ttataaaccc
30781 30841 30901 30961	gtacccacaa cttcaaccct agggtttatt tttaaccca	tcttcatgtc gtctacccct tccccaaatg ctaacaacca	atgaagatga gcttcacaca caggcggatc	aagcacctcc aagcccagac tctacagcta	caacacccct ggagttctta aaagtgggag	ttataaaccc ctttaaaatg ggggacttac
30781 30841 30901 30961	gtacccacaa cttcaaccct agggtttatt tttaaccca	tcttcatgtc gtctacccct tccccaaatg ctaacaacca	atgaagatga gcttcacaca caggcggatc	aagcacctcc aagcccagac tctacagcta	caacacccct ggagttctta aaagtgggag	ttataaaccc ctttaaaatg ggggacttac
30781 30841 30901 30961 31021	gtacccacaa cttcaaccct agggtttatt tttaacccca agtggatgac	tcttcatgtc gtctacccct tccccaaatg ctaacaacca actgatggta	atgaagatga gcttcacaca caggcggatc ccttacaaga	aagcacctcc aagcccagac tctacagcta aaacatacgt	caacacccct ggagttctta aaagtgggag gctacagcac	ttataaaccc ctttaaaatg ggggacttac ccattactaa
30781 30841 30901 30961 31021 31081	gtacccacaa cttcaaccct agggtttatt tttaacccca agtggatgac aaataatcac	tcttcatgtc gtctacccct tccccaaatg ctaacaacca actgatggta tctgtagaac	atgaagatga gcttcacaca caggcggatc ccttacaaga tatccattgg	aagcacctcc aagcccagac tctacagcta aaacatacgt aaatggatta	caacacccct ggagttctta aaagtgggag gctacagcac gaaactcaaa	ttataaaccc ctttaaaatg ggggacttac ccattactaa acaataaact
30781 30841 30901 30961 31021 31081	gtacccacaa cttcaaccct agggtttatt tttaacccca agtggatgac	tcttcatgtc gtctacccct tccccaaatg ctaacaacca actgatggta tctgtagaac	atgaagatga gcttcacaca caggcggatc ccttacaaga tatccattgg	aagcacctcc aagcccagac tctacagcta aaacatacgt aaatggatta	caacacccct ggagttctta aaagtgggag gctacagcac gaaactcaaa	ttataaaccc ctttaaaatg ggggacttac ccattactaa acaataaact

FIG. 28A-8

31201	tattaacacc	ttatggactg	gaataaaccc	tccacctaac	tgtcaaattg	tggaaaacac
31261	taatacaaat	gatggcaaac	ttactttagt	attagtaaaa	aacggagggc	ttgttaatgg
21221	ctacatatat	ctacttcctc	tatcagacac	totoaaccaa	atgttcacac	aaaagacagc
21201	aaacatccaa	ttaarattat	attttgactc	ttctggaaat	ctattaactg	atgaatcaga
31301	cttaaaaatt	ccacttaaaa	ataaatcttc	tacagegace	agtgaaactg	tagccagcag
31601	caaagccttt	ataccaaata	ctacagetta	tcccttcaac	accactacta	gggatagtga
31301	aaactacatt	catogaatat	ottactacat	gactagttat	gatagaagtc	tatttccctt
3120T	gaacatttct	atastactas	acaccotat	gatttcttcc	aatgttgcct	atoccataca
31021	atttgaatgg	acaatgecaa	acageegeae	tecagaaage	aacatagcta	cactaaccac
3108T	atccccttt	ttatttatt	acattacaca	aracracaac	taaaataaag	tttaagtgtt
31/41	tttatttaaa	ateressant	togactage	attttacctc	caccttccca	tttgacagaa
3180T	tacaccaatc	tetacaaaa	cacacettta	aacatttooa	taccattaga	gatagacatt
3180T	gttttagatt	coccccacg	aacagtttca	dacaccaga	atctggggtc	agtgatagat
31921	aaaaatccat	ccacatteca	ttttaaaaaa	ctttcacant	ccaactocto	cogatocgaa
31381	tccggagtct	egegacagee	astataasaa	aaraarratr	ngaatcataa	tccgaaaacg
32041	gtatcggacg	ggatcacggt	catcuygaay	aagaacgacg	ctatctacat	cactccatac
32101	aactgctgtt	attgtgtete	accadaccca	tatactasa	catcatttta	atagecetta
32161	aactgctgtt	tatgggatea	because	aaccettct	gatttcactc	asatctttgc
32221	acatcaactt	tctggtgcga	egegegeage	ttaataaaa	staattaaaa	gegeterage
32281	agtaggtaca	acacattatt	acaatattgt	characata	ataccasact	ttaatataaa
32341	caaaactcat	atctgatata	ategeeetg	catgattatt	ataccaaage	caatataca
32401	ttaaatgacg	ttccctcaaa	aacacactac	ccacatacat	gatetette	astataacct
32461	tattaacaat.	ctgtctgtac	catggacaac	gttggttaat	catgeaacce	gactactact
32521	tccggaacca	cactgccaac	accgctcccc	cagecatgea	ccgaagcgaa	tennanatat
32581	tacaatgaca	atgaagaacc	caattctctc	gaccgtgaat	cacttgagaa	cyaaaaaaaa
32641	ctatagtggc	acaacataga	cataaatgca	tgcatcttct	cataattttt	addiction
32701	gatttagaaa	catatcccag	ggaataggaa	gctcttgcag	aacagtaaag	ctygeagaac
32761	aaggaagacc	acgaacacaa	cttacactat	gcatagtcat	agtatcacaa	tetggeaaca
32821	gegggtggte	ttcagtcata	gaagctcggg	tttcattttc	ctcacaacgt	ggtaactggg
22991	ctctaatata	agggtgatgt	ctaacacata	atotcoacco	tgcgcgcaac	cttgtcataa
329/1	tagaattact	tcctgacatt	ctcgtatttt	gtatagcaaa	acgcggccct	ggcagaacac
33001	actettette	geettetate	ctgccgctta	gcgtgttccg	tgtgatagtt	caagtacage
33061	cacactetta	agttggtcaa	aagaatgctg	gcttcagttg	taatcaaaac	tecategeat
22121	ctaattotto	toaooaaatc	atccacggta	gcatatgcaa	atcccaacca	agcaatgcaa
33181	ctggattgcg	tttcaagcag	gagaggagag	ggaagagacg	gaagaaccat	gctaattttt
33241	attccaaacq	atctcgcagt	acttcaaatt	gtagatcgcg	cagatggcat	ctctcgcccc
33301	cactgtgttg	gtgaaaaagc	acagctaaat	caaaagaaat	gcgattttca	aggtgctcaa
22261	caataacttc	caacaaagcc	tccacgcgca	catccaagaa	caaaagaata	ccaaaagaag
33421	gaggattttc	taactcctca	atcatcatat	tacattcctg	caccattccc	agataatttt
22/21	cacctttcca	geettgaatt	attcototca	attettataa	taaatccaat	ccacacatta
33541	caaacaggtc	ccaaaaaaca	ccctccacca	ccattcttaa	acacaccctc	ataatgacaa
33601	aatatettee	tectatatea	cctataacaa	attgagaatg	gcaacatcaa	ttgacatgcc
33661	cttggctcta	agttettett	taagttctag	ttgtaaaaac	tctctcatat	tatcaccaaa
33721	ctdcttadcc	agaagccccc	coogaacaag	agcaggggac	gctacagtgc	agtacaagcg
33721	cagacctccc	caattoocto	caccaaaaac	aagattggaa	taagcatatt	gggaaccgcc
220/1	agtaatatca	traaanttar	togaaatata	atcaggcaga	gtttcttgta	aaaattgaat
33041	2222722222	tttaccasas	aaacattcaa	aacctctggg	atocaaatoc	aataggttac
22061	cacactacaa	treateatte	ttagttttga	attagtctgc	aaaaataaaa	aaaaaaacaa
33301	cgcgccgcgc	atactacct	racraacard	tggataaatg	agtettteca	tcacaagaca
34021	gegeeacace	totocageee	gacgaccagg	aaacctotca	tootoattaa	acaacagcac
34081	agecacaggg	tagagatasa	gaccetegea	aattetteat	gaagcataca	atccagacat
34141	cyaaaguuco	-ttp://grad	cagcacgaac	aacatagcct	ttoootataa	ttatgcttaa
3420T	gttagcatca	gilaacyaya	aaaaacagcc	atacaaaacta	aaaggcacac	gagaataaaa
34261	tegtaagtat	aycaaaycca	- ccccccgcgg	caacatagaa	cccatccct	ctaaatacac
34321	aacacaacca	tttetetget	gergereagg	gagagagtag	. cccggcccc	cacaanctct
34381	atacaaagco	catcagcca	Lygoriacia	tatacacayid	. agogggcacg	cacaagctct
34441	aaagtcactc	ccaacctct	ccacaatata	-acadacadg	aree at are	acgtaatggg
34501	agtaaagtgt	. aaaaaatccc	gccaaaccca	acacacaccc	togadautgug	tcaccaggga
34561	aaagtacagt	. ttcacttccg	caatcccaac	aagcgtcact	. Lectettet	cacggtacgt
34621	. cacatcccat	: taacttgcaa	cgtcattttc	ccacggccgc	geegeeegt	ttagccgtta
34681	. accccacago	: caatcaccac	: acaccccaca	attttaaaa	tcacctcatt	tacatattgg
34741	caccattcca	ı tctataaggt	. atattattga	. tgatg		
SEQ I	D NO: 12					

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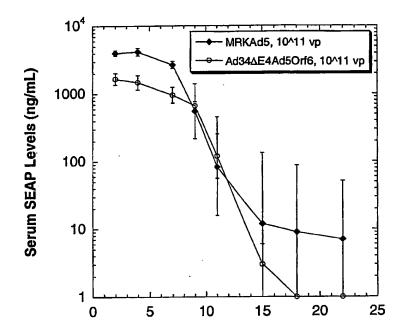


FIG. 29

Vaccine	Monkey	Р	re	W	k 4	W	k 8	W	24	W	28	¥	36
Wk 0, 4, 24	ID _	Mock	Gag*	Mock	Gag	Mack	Gag	Mock	Gag	Mock	Gag	Mock	639
MRKAd5gag, 1041 vp MRKAd5gag, 1041 vp MRKAd5gag, 1041 vp	00C018 00C034 00C058	1 0 4	5 4 4	13 5 3	1025 219 1086	0 5 0	824 404 440	8 3 4	756 445 1439	D 3 0	474 339 2338	0 0 0	383 216 940
Ad34AE1gagAE4Ad5Orf6, 10*11 vp Ad34AE1gagAE4Ad5Orf6, 10*11 vp Ad34AE1gagAE4Ad5Orf6, 10*11 vp	00D038 00D042 00D066	6 6 3	8 30 18	5 4 1	111 89 118	1 4 1	301 264 816	0 1 0	224 73 429	1 0 0	535 181 439	0 0	233 69 273

Vaccine	Monk ID	•	D4 <sup>+</sup> CD3 <sup>+</sup> mphocytes	IFN-γ <sup>+</sup> CD8 <sup>+</sup> CD3 <sup>+</sup> per 10 <sup>6</sup> Lymphocytes		
		Mock	Gag <sup>a</sup>	Mock	Gag <sup>a</sup>	
Ad34ΔE1gagΔE4Ad5Orf6	00D038	22	154	130	450	
	00D042	32	118	96	171	
•	00D066	12	238	150	442	

Vaccine	vaccine Vaccine		P	re	T=4	wks	T=8 wks		Ts24 wits		T=26 wks		T=32 wks	
T=0, 4 wks	T=24 wks	ED.	Mock	Cing*	Mock	Gag	Mock	Cag	Mock	ď	Mock	Gag	Mock	Chag
Ad345E1gag6E4Ad5Od6, 10*11 vp	Ad35&E1gag&E4Ad5Orf8, 10^10 vp	000016	4	8	٦.	84	5	334	5	29	0	305	3	244
Ad34AE1gag&E4Ad3Orf8, 10*11 vp	Ad35AE1gagAE4Ad5Orf8, 10^10 vp	000044	1	1	8	79	0	374	В	138	0	493	1	253
Ad346E1gzg6E4Ad5Orf6, 10^11 vp	Ad35&E1gag&E4Ad5Orl6, 10*10 vp	00D064	4	8	1	125	8	655	6	145	٥	351	1	235
	<del> </del>		-	<del></del> -	<b>—</b>	<del></del>	-	54	٠.	<del></del> -	-	<u> </u>		<u> </u>
NaNa		00D087			<u> 3</u> _	3	_ 8	_ 54	<u> </u>			<u> </u>		<u> </u>

Vaccine (Y=0, 4 Wks)	Vaccine (T=24 Wk)	Monkey		D4*CD3* mphocytes	IFN-y*CD8*CD3* per 10 <sup>8</sup> Lymphocytes		
		ID	Mock	Gag	Mock	Gag	
Ad34AE1gagAE4Ad5Orf6, 10^11 vp	Ad35ΔE1gagΔE4Ad5Orf6, 10^10 vp	00D016	62	433	176	1288	
Ad34∆E1gag∆E4Ad5Orf6, 10^11 vp	Ad35∆E1gag∆E4Ad5Orf6, 10^10 vp	00D044	136	593	323	1871	
Ad34ΔE1gagΔE4Ad5Orf6, 10^11 vp	Ad35ΔE1gagΔE4Ad5Orf6, 10^10 vp	00D064	188	785	292	892	

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